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RESULT 79
ADA26875/
ID ADA2
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AC ADA2
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DT 20-N
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DB Huma
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KW Meta
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 multiplex PCR amplification, for genetic testing or facilitating multiplex PCR amplification from limiting amounts of target nucleic acid. The methods are also useful for improving genetic diagnostic and screening methods, such as prenatal diagnostic testing, foetal sex determination or genetic identification, e.g. DNA profiling or DNA fingerprinting. The nucleic acid sequence amplification is also useful in forensic analysis of degraded, old, anotient and difficult samples that are difficult to amplify and identify. This sequence represents a PCR primer used in the selection and amplification of genetic markers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Selecting genetic markers as targets for nucleic acid sequence amplification, useful for improving genetic testing, e.g. fetal determination, comprises selecting each of the genetic markers at a heterozygosity index.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genetic marker selection; multiplex PCR amplification; prenatal diagnostic testing; foetal sex determination; genetic identification; DNA profilling; DNA fingerprinting; forensic analysis; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention describes a method of selecting genetic markers as targets for nucleic acid sequence amplification comprising selecting each of the genetic markers according to a heterozygosity index of 0.5 or greater. Selecting and amplification of genetic markers are useful as targets for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003031646-A1
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                                                                               gastrointestinal; prostate; breast; colorectal; diagnostic imaging;
drug targeting; chromosome 8q24.3; human;
protein tyrosine phosphatase type IVA member 3; PRL-3; gene mapping;
                                                                                                                                                                                                                                                                                                                                                                                                          ADA26875
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12-OCT-2001;
                                                                                                                                                                                                                                              Human PRL-3
                                                                                                                                                                                                                                                                                                 20-NOV-2003
  Homo sapiens
                                                        cytostatic;
                                                                                                                                                                                        Metastasis; neoplastic
                                                                                                                                                                 neoplastic
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19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                          standard;
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                                                                                                                                                         ; neoplastic growth; detection; growth marker; drug screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matthews
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                                                                                                                                                                                                                                           forward PCR primer #159, used in gene mapping
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2001AU-00008235
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                                                        primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                 entry)
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95.0%;
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                                                                                                                                                              owth; detection; prediction; drug screening; cancer; tumour;
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Pred. No. 1.3e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 Other;
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TXAXEXEXEXEXE UXAXEXEXEXEXE

Intestinal 20-MAY-2004

(first entry)

inflammatory

intestinal epithelium cell development; peyer's patch M cell development;

epithelium/peyer's patch M cell-associated PCR primer #93.

ry bowel disease; glutenenteropathy; infectious disease; haemolytic anaemia; rheumatoid arthri

arthritis;

닭 S

20

ADL24948 standard;

DNA;

20

Matches Query Match Best Local 9

19;

Conservative

0

Mismatches

No. 1

.3e+03;

DB 1;

Length Indels

0

Gaps

0

Similarity

1.9%; 95.0%;

Score 18.4; Pred.

374 CTGCCTCAGCCTCCCAAAGT 393

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The invention relates to methods for identifying regions of neoplastic components of metastasis. The methods involve determining whether a neoplastic growth care protein is overexpressed, either by the use of an antibody committee protein is overexpressed, either by the use of an antibody component committee a disclosed (SAGE (serial analysis of gene expression) tags for these are given in ADA26759-ADA26796), with protein tyrosine phosphatase type IVA member 3 (also known as pRL-3) being a preferred neoplastic growth markers are disclosed (SAGE (serial analysis of gene expression) tags for type IVA member 3 (also known as pRL-3) being a preferred neoplastic growth markers are specifically expressed at a higher level in metastatic growth markers are specifically expressed coverexpression of the neoplastic growth markers are specifically expressed that the tissue has a propensity to metastasis. The invention also concompasses methods for treating a patient with an advanced or metastatic cancers. The methods of the invention are useful for identifying regions of neoplastic growth, for detecting or predicting metastasis. Or identifying candidate drugs for treating advanced or metastatic cancers. The methods of the invention are useful for gastrointestinia, prostate, breast or colorectal cancers. Antibodies which bind to the neoplastic growth marker proteins are additionally useful for diagnostic imaging and for targeting cytotoxic or chemotherapeutic drugs. The present sequence represents a PCR primer used concents and the PRF.-1 reneared med and a propensity and for targeting cytotoxic or metastaric cancers. The invention are december appreciation are useful for the neoplastic growth marker proteins are additionally useful for diagnostic imaging and for targeting cytotoxic or chemotherapeutic drugs. The present sequence represents a PCR primer used to chemotherapeutic drugs. The present sequence are presents a protein and primer used to chemotherapeutic drugs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying regions of neoplastic growth in a human body, useful for detecting or predicting metastasis, comprises administering to the human body an antibody or peptide that specifically binds to a protein marker
                                        chemotherapeutic drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-OCT-2002;
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Sequence
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BP;
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       0 Other;
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RESULT 795
ADL25083
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XX ADL250
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XX Inte
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention comprises DNA sequences which are associated with intestinal epithelium and peyer's patch M cells. The DNA sequences of the invention are useful for assessing, modifying, modulating or regulating intestinal epithelium or M cell development. The DNA sequences of the invention are also useful in the treatment of: inflammatory bowel disease, glutementeropathy, infectious disease, autoimmune diseases (e.g. haemolytic anaemia, rheumatoid arthritis, derawe's disease, multiple sclerosis, allergy, asthma and diabetic mellitus), diseases or disorders of the immune system, hypersensitivity, anaphylaxis, and blood group incompatibility. The present DNA sequence represents a PCR primer that was used to amplify an intestinal epithelium/peyer's patch M cell-associated DNA sequence of the invention.
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      04-APR-2002; 2002WO-US010873.
                                                                                                                                                                                                                                      Grave's disease; multiple sclerosis; allergy; asthma; immune system disorder; hypersensitivity; anaphylaxis; blood group incompatibility; ss; human; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel isolated or purified polypeptide encoded by genes associated with intestinal epithelium or M cell development, differentiation or function useful for treating autoimmune diseases and infectious diseases.
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                                                                 17-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADL25083 standard;
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                                                                                                                            WO200280852-A2
                                                                                                                                                                                                                                                                                                                             intestinal epithelium cell development; peyer's patch M cell development; inflammatory bowel disease; glutenenteropathy; infectious disease; autoimmune disease; haemolytic anaemia; rheumatoid arthritis; dermatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Intestinal epithelium/peyer's patch M cell-associated PCR primer #228.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                  intestinal epithelium cell development;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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Pred. No. 1
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                                                                                                                                                                                                                                                                        anaphylaxis;
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                                                                                                                                                                                                                                                                                                       diabetic
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RESULT 796
ABD30939
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                            Nyce JW,
Miller S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                intestinal epithelium and peyer's patch M cells. The DNA sequences of the invention are useful for assessing, modifying, modulating or regulating intestinal epithelium or M cell development. The DNA sequences of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 593; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel isolated or purified polypeptide encoded by genes associated with intestinal epithelium or M cell development, differentiation or function
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Brayden DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-APR-2001; 2001US-0281416P
                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         emphysema;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human RANTES-derived oligonucleotide SEQ ID 13150.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABD30939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-075470/07
                                                                                                                                                                                                                         23-APR-2002; 2002WO-US013143
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                                                                                                               (EPIG-)
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                                                                                                               EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
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                               Li Y,
Tang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGCGATCCTCCTGTCTCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                               chronic obstructive pulmonary disease; transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Byrne D,
                                                                                                                                                                     2001US-0286036P
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                               Ļ
                               Sandrasagra A,
L, Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bronchoconstriction; allergy; hyposecretion; pain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 18.4;
Pred. No. 1.
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                                                              Pabalan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   are associated with
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                         Aguilar
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cysturction or cancer and can be anti-recibe to the contresponding mayn.

CC The invention also describes a kit, that comprises: (a) a delivery

CC device, in separate containers, (b) the oligonucleotides, (c)

co fine invention for adding a carrier and for use of the kit. The composition

CC of the invention has antiallergic, antiinflammatory, antiasthmatic,

CC analysesic, hypotensive, immunosuppressive and cytostatic activity, is a

CC treating a respiratory, lung or malignant disease. The administered

CC composition comprises oligo and is administered to reduce the production

CC or availability, or to increase the degradation of the target mRNA or to

CC reduce the amount of target polypeptide present in the lungs. The

CC inflammation, allergies and/or bronchoconstriction and/or lung

CC with a disease or condition such as pulmonary vasoconstriction

CC inflammation, allergies, asthma, impeded respiration, respiratory

CC distress syndrome, pain, cystic fibrosis, allergic rithinitis, pulmonary

CC inflammation, rejection, pulmonary infections, bronchitis or cancer.

CC The reduced adenosine content of the anti-sense oligos corresponding to

CC thymidines present in the target RNA serves to prevent the breakdown of

CC the oligonucleotides into products that free adenosine into the system

CC prevent any unwarted affect disease, tissue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  밁
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Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        comprising oligonucleotides, effective for alleviating
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                                                                                                                                                                                         Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hyportension; respiratory distress syndrome; allergic rhinitis; pulmonary hyportension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bronchoconstriction, respiratory tract inflammation, allerg reducing adenosine sensitivity, levels of adenosine (A) or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABD31043 standard; DNA; 20
                                                                    Homo sapiens.
                                                                                                                                      emphysema; chronic obstructive pulmonary
pulmonary transplantation rejection; ss;
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Pred. No. 1.
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                                                                                                                                      primer.
                                                                                                                                                                        disease; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
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WO200285309-A2

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29-JUL-2004 ABD32136;

(first entry)

Human PDE4C-derived oligonucleotide SEQ ID

RESULT 798

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ABD32136

ABD32136 standard; DNA; 20

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cc surfactant depletion or hyposecretion, when administered to a mammal. The capture of dysfunction of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, cc analgesic, hypotensive, immunosuppressive and cytostatic activity, is a cc beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered cc composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The creduce the amount of target polypeptide present in the lungs. The cultinonary obstruction, and/or bronchoconstriction and/or lung crifilammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, comprises syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary crification, pulmonary conditions, bronchoconstriction, pulmonary conditions, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system content on the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the reby, to or prevent any unwarted affects due to it
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                                                                   Query Match
Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allers reducing adenosine sensitivity, levels of adenosine (A) or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-APR-2001; 2001US-0286036P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes a novel composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    bronchodilating agent.
                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (EPIG-) EPIGENESIS PHARM INC
                                 636 TCTGTCACCCAGGCTGGAGT 655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15; SEQ ID NO 13254; 763pp;
                                                                       19;
                                                                                      Similarity
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, Tang L,
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                                                                                                                                             B₽;
                                                                                                                                                                                  unwanted effects due
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L, Shahabuddin
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                                                                                                                                             C; 7
                                                                       0
                                                                                                                                               G; 5 T; 0 U; 0 Other;
                                                                                        Score 18.4;
Pred. No. 1
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surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung alray or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA.

The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cycostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or surfactant hymogradiation and/or lung.
                                                                                                               Query Match
Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                             inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                             Sequence
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CTGGTCTCAAACTCCTGACC 1134
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                                                                                                                   Conservative
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L, Shahabuddin
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CC comprising oligonucleotides, effective for alleviating
CC control of the composite sensitivity, levels of adenosine (A) or (A) receptors,
CC surfactant depletion or hyposecretion, when administered to a mammal. The
CC oligonucleotides are derived from a gene encoding or regulating
CC expression of a target polypeptide associated with lung airway or lung
CC dysfunction or cancer and can be anti-sense to the corresponding mRNA.
CC The invention also describes a kit, that comprises: (a) a delivery
CC device, in separate containers, (b) the oligonucleotides, (c)
CC instructions for adding a carrier and for use of the kit. The composition
CC of the invention has antiallergic, antiinflammatory, antiasthmatic,
CC analgesic, hypotensive, immunosuppressive and cyrostatic activity, is a
CC beta-adrenergic agonist. The composition is useful for preventing or
CC composition comprises oligo and is administered to reduce the production
CC or availability, or to increase the degradation of the target mRNA or to
CC reduce the amount of target polypeptide present in the lungs. The
CC inflammation, allergies and/or bronchoconstriction and/or lung
CC with a disease or condition such as pulmonary vasoconstriction,
CC inflammation, allergies, asthma, impeded respiration, respiratory
CC distress syndrome, panin, cystic fibrosis, allergic rhinitis, pulmonary
CC transplantation rejection, pulmonary infections, bronchitis or cancer.
CC manifestion the composition of the target pulmonary
CC manifestion rejection, pulmonary infections, bronchitis or cancer.
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Miller S
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transplantation rejection, pulmonary infections, bronchitis or cancer The reduced adenosine content of the anti-sense oligos corresponding thymidines present in the target RNA serves to prevent the breakdown the oligonucleotides into products that free adenosine into the syste
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 15; SEQ ID NO 13255; 763pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acids associated with lung airway or lung dysfunction, and
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CC comprising oligonucleotides, effective for alleviating controlled the producting adenostriction, respiratory tract inflammation, allergies and CC reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, CC surfactant depletion or hyposecretion, when administered to a mammal. The CC oligonucleotides are derived from a gene encoding or regulating CC expression of a target polypeptide associated with lung airway or lung CC dysfunction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production
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                                                                                                                                                                                                                                                                                                                Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targenucleic acids associated with lung airway or lung dysfunction
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                                                                                                                                                                                                                                                                                                   bronchodilating agent.
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or availability, or to increase the degradation of the target reduce the amount of target polypeptide present in the lungs pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABD30933 standard; DNA; 20 BP
                                                                                                                Pharmaceutical composition oligonucleotide containing nucleic acids associated wi
                                                                                                                                                                          WPI;
                                                                                                                                                                                                       Miller
                                                                                                                                                                                                                                                                              24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                           23-APR-2002; 2002WO-US013143
                                                                                                                                                                                                                                                                                                                                                                     WO200285309-A2
                                                                                                                                                                                                                                                                                                                                                                                                  Homo
                                                                                                                                                                                                                                                                                                                                                                                                                              pulmonary transplantation
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                                                                                                                                                                                                                                                  (EPIG-)
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                                                                                                                                                                                                      Li Y, Sa
Tang L,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                       Sandrasagra A,
L, Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oligonucleotide SEQ ID 13144.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                               rejection; ss; primer.
                                                                                                              for treating asthma, has antisense less percentage of adenosine, targeted to ith lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 G; 4 T; 0 U;
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                                                                                                                                                                                                                        Katz
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No. 1.3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 Other;
                                                                                                                                                                                                                        Pabalan
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comprising oligonucleotides, bronchoconstriction, respirat

cleotides, effective n, respiratory tract sensitivity, levels

sition (a) a first active for alleviating inflammation, allergies a of adenosine (A) or (A) r

receptors

agent,

describes a novel composition

Claim

15;

SEQ

IJ

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13144;

763pp;

English

bronchodilating agent

with

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RESULT 802
ABD26091/c
ID ABD260
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Best Local S
Matches 19
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      Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; antisense; bronchoconstriction, allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
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                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AA463249-derived
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                                                                                                                                                               24-APR-2001; 2001US-0286036P
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                                                                                                    EPIGENESIS
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Li Y,
Tang
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotide SEQ ID 5103
      Sandrasagra A,
L, Shahabuddin
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                                                                                                    PHARM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 18.4; DB 1; Pred. No. 1.3e+03;
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                                       Katz
                                   'n
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                                       Pabalan
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                                   Aguilar
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comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating control adenosine sensitivity, levels of adenosine (A) or (A) receptors, consider and expectant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating control or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery construction for adding a carrier and for use of the invention has antiallergic, antiinflammatory, antiasthmatic, analyssic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or reduce the amount of target polypeptide present in the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, allergies and/or bronchoconstriction are associated with a disease or condition such as pulmonary vasoconstriction are associated with a disease or condition such as pulmonary vasoconstriction are associated with a disease or condition such as pulmonary vasoconstriction are associated with a disease or condition such as pulmonary vasoconstriction are associated with a disease or condition such as pulmonary vasoconstriction are associated with a disease or condition such as pulmonary vasoconstriction are associated conformation, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system c.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                bronchodilating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15; SEQ ID NO 5103;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               describes
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Query Match Best Local Similarity Sequence 20 BP; 4 A; 4 C; 9 G; 3 T; 0 U; 0 Other; Score 18.4; Pred. No. 1 DB 1; Length 20;

밁 S 970 20 TCGGCTCACTGCAACCTCCG 1 TCGGCTCACTGCAACCTCTG 989

Matches

19;

Conservative

o ;:

Mismatches

1:

Indels

0

Gaps

0

1.3e+03

95.0%;

ABD26094 standard; DNA; 20

AA463249-derived oligonucleotide SEQ ID 5106.

29-JUL-2004 ABD26094;

(first entry)

RESULT 803
ABDZ6094/c
ID ABDZ60
XX Y
XX Human;
KW respir
KW surfac
XX surfac
XW surfac
XW surfac
XW emphys
EW beta-a
KW respir
KW respir
KW beta-a
KW pulmon
XX Pu WO2002
XX emphysema; pulmonary t Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; transplantation

Homo sapiens.

WO200285309-A2

WPI; 2003-093058/08

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ABD30934
ID ABD
XX
AC ABD
XX
DT 29-
XX
DE Hum
XX
KW Hum
                                                                                                                                                                                                                                                                                                                                                                                                                   CC surfactant depletion or hyposecretion, when administered to a mammal. The copression of a target polypeptide associated with lung airway or lung CC dysfunction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered CC composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The CC pulmonary obstruction, and/or bronchoconstriction and/or lung conformation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, comparises, asthma, impeded respiration, respiratory conformation, emphysema, chronic obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary the reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system content on the carget RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the respective pulmonary and effects due to it.
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                                                                                                                                                                            RESULT 804
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) recurred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targ nucleic acids associated with lung airway or lung dysfunction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 15; SEQ ID NO 5106;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             bronchodilating
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   Human;
                                     Human RANTES-derived oligonucleotide SEQ ID 13145
                                                                        29-JUL-2004
                                                                                                                                           ABD30934 standard; DNA;
                                                                                                                                                                                                                                                                    478
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19; Conser
   antisense; bronchoconstriction; allergy; hyposecretion; pain;
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Tang L,
                                                                                                                                                                                                                                                                    AAGTGCAGTGGTGATCAC 497
                                                                                                                                                                                                                                    AAGTGCAGTGGTGATCTC 1
                                                                                                                                                                                                                                                                                                                                                                        BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                             unwanted effects due to it
                                                                                                                                                                                                                                                                                                       Conservative
                                                                        (first entry)
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L, Shahabuddin
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                        1.3e+03
                                                                                                                                                                                                                                                                                                                                                                             0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              adenosine, targeted to lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                        Length 20
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CC comprising oligonuclectides, effective for alleviating allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, comprising of a target polypeptide associated with lung airway or lung compression of a target polypeptide associated with lung airway or lung compression of a target polypeptide associated with lung airway or lung compression of a target polypeptide associated with lung airway or lung compression of a target polypeptide associated with lung airway or lung compression of a target polypeptide associated with lung airway or lung compression of a target polypeptide associated with lung airway or lung compression and for use of the corresponding mRNA.

CC dysingtion or cancer and can be anti-sense to the corresponding mRNA.

CC device, in separate containers, (b) the oligonuclectides, (c) analgesic, hypotensive, immunosuppressive and cytostatic activity, is a composition has antiallergic, antiinflammatory, antiasthmatic, compressive, immunosuppressive and cytostatic activity, is a composition comprises oligo and is administered to reduce the production composition and administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition such as pulmonary vascoconstriction, allergies and/or bronchoconstriction and/or lung inflammation, allergies asthma, impeded respiration, respiratory condition such as pulmonary vascoconstriction, respiratory conditions with a such as pulmonary disease, pulmonary disease, pulmonary conditions, pulmonary disease, pulmonary conditions, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the noile pulmonary diseases.
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oligonucleotides into products that free adenosine into the system , lung, brain, heart, kidney, etc. tismus and the system vent any uncorrect.
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           The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, allergies and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system of ce... lung, brain, heart, kidney, etc, tissue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respirator, tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) recepto surfactant depletion or hyposecretion, when administered to a mammal oligonucleotides are derived from a gene encoding or regulating
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Tang L,
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        CC composition of igonucleotides, effective for alleviating controlled from the controlled from a gene encoding or (A) receptors, cc surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating cc expression of a target polypeptide associated with lung airway or lung cdysfunction or cancer and can be anti-sense to the corresponding mRNA. cc The invention also describes a kit, that comprises: (a) a delivery cc device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, cc analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or activative or the invention of the invention of the invention of the invention of the composition of the invention of the inventio
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
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                                                                                                                                                                                                                    Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human PDE4C-derived oligonucleotide SEQ ID 14304.
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comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The
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                                                                                                   This invention describes a novel composition
                                                                                                                                                    Claim 15; SEQ ID NO 14304;
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Shahabuddin
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Pred. No. 1.3e+03;
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cc oligonucleotides are derived from a gene encoding or regulating cc expression of a target polypeptide associated with lung airway or lung cdysfunction or cancer and can be anti-sense to the corresponding mRNA. Cc The invention also describes a kit, that comprises: (a) a delivery cd device, in separate containers, (b) the oligonucleotides, (c) constructions for adding a carrier and for use of the kit. The composition cc of the invention has antiallergic, antiinflammatory, antiasthmatic, and creating a respiratory, lung or malignant disease. The administered composition composition composition is useful for preventing or creating a respiratory, lung or malignant disease. The administered correspondint, or to increase the degradation of the target mRNA or to creduce the amount of target polypeptide present in the lungs. The cultiform or availability, or to increase the degradation of the target mRNA or to creduce the amount of target polypeptide present in the lungs. The cultiform such as pulmonary vasoconstriction are associated with a disease or condition such as pulmonary vasoconstriction are associated with a disease or condition such as pulmonary vasoconstriction are associated with a disease or condition such as pulmonary vasoconstriction are pulmonary condition, allergies, asthma, impeded respiration, respiratory conditions, allergies, asthma, impeded respiration, respiratory conditions, pronchitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the production and for the aritem or the artem of the artem of the interesponding to the oligon corresponding to the production of the artem of the prevent the breakdown of the production of the artem of the p
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; antigesic; hypotensive; immunosuppressive; cytostatic; cytic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction respiratory distress syndrome; allergic rhinitis; pulmonary hypertension respiratory distress syndrome; allergic rhinitis; pulmonary hypertension
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                                                           Li Y, Sa.
Tang L,
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                                                                      Sandrasagra A,
L, Shahabuddin
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Pred. No. 1.3e+03;
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                                                                                                                                                                               Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension;
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                                                                                                                                                                                                                                                                                                                                               Human RANTES-derived oligonucleotide SEQ ID 13206.
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                                                                                                                                      emphysema; chronic obstructive pulmonary disease; cancer;
pulmonary transplantation rejection; ss; primer.
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Pred. No. 1.
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SAXBXBXXX

standard; DNA;

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Human; antisense;

bronchoconstriction; allergy; hyposecretion; pain; inflammation; adenosine sensitivity; lung; cancer;

Human RANTES-derived oligonucleotide SEQ ID 13245.

(first entry)

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                                                                                                                                                                   cc analgesic, hypotensive, immunosuppressive and cytostatic activity, is a cc beta-adrenergic agonist. The composition is useful for preventing or cc treating a respiratory, lung or malignant disease. The administered cc composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to creduce the amount of target polypeptide present in the lungs. The cc pulmonary obstruction, and/or bronchoconstriction and/or lung ulmonary obstruction, and/or surfactant hypoproduction are associated cc with a disease or condition such as pulmonary vasoconstriction, cc inflammation, allergies and/or surfactant hypoproduction are associated cc with a disease or condition such as pulmonary vasoconstriction, cc inflammation, allergies, asthma, impeded respiration, respiratory cc distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary cc transplantation rejection, pulmonary infections, bronchitis or cancer. cc The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system cc e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwarted affect disc.
                                                                               Query Match
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Matches 19
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                                                                                                                                                                   Sequence
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                                                                                                     Similarity
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                                                                                                   1.9%;
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                                                                                                     Score 18.4;
Pred. No. 1.
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                                                                                    Mismatches
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                                                                                                 1.3e+03;
                                                                                                                         DB 1;
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                                                                                                                       Length 20
                                                                                    Indels
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canalgesic, hypotensive, immunosuppressive and cyrostatic activity, is a contract advanced advanced against. The composition is useful for preventing or composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to creduce the amount of target polypeptide present in the lungs. The complementary obstruction, and/or bronchoconstriction and/or lung confinamation, allergies and/or surfactant hypoproduction are associated confinamation, allergies and/or surfactant hypoproduction are associated confinamation, allergies, asthma, impeded respiration, respiratory confinamation, emphysema, chronic obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary confiness syndrome pain, cystic fibrosis, allergic rithinitis, pulmonary confiness syndrome pain, cystic fibrosis, allergic rithinitis, pulmonary confiness present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system content on the serves to pracent the breakdown of the oligonucleotides into products that free adenosine into the system content any immanted affects dise to it.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           reducing adenosine sensitivity, levels of administered to a mammal. The surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, is a containers immunosuppressive and cytostatic activity, is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nyce JW,
Miller S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors.
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Sequence
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Tang
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    BP; 5
                                                                                      unwanted effects
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L, Shahabuddin
P
4 C; 8
G; 3 T; 0 U; 0 Other;
                                                                                      due to it
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tis; pulmonary hypertension;
ase; cancer; bronchitis;
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(EPIG-)

EPIGENESIS PHARM INC

WAN AND SOLUTION OF THE STATE O

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RESULT 811
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; antisense; bronchoconstriction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABD30940;
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pain;

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and Nyce JW, Miller S WPI; 2003-093058/08 ß Li Y, Tang Sandrasagra A, L, Shahabuddin ŝ

Katz 'n

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Aguilar

SEQ ID NO 13151; 763pp; English

agent

canalgesic, hypotensive, immunosuppressive and cytostatic activity, is a compelse composition of useful for preventing or composition comprises oligo and is administered to reduce the production comprises oligo and is administered to reduce the production of the target polypeptide present in the lungs. The complomary obstruction, and/or bronchoconstriction and/or lung confilmmation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasconstriction, and/or surfactant hypoproduction are associated confilmmation, allergies, asthma, impeded respiration, respiratory confilmmation, allergies, asthma, impeded respiration, respiratory confirmmation, allergies, asthma, impeded respiration, respiratory confirmation respection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to configurate present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system confirmation in the product of the configuration in the street what free adenosine into the system confirmation in the products that free adenosine into the system confirmation in the product of the configuration of the configuration in the product of the configuration of the configur comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, is a second of the invention of the invention has antiallergic, antiinflammatory, antiasthmatic, This invention describes a novel composition (a) a first active agent,

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mana or to reduce the amount of target polypeptide present in the lungs. The
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 15;
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Tang
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, Shahabuddin
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RESULT 813
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nyce JW,
Miller S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antisthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-APR-2001; 2001US-0286036P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N58473-derived oligonucleotide SEQ ID 7966.
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19; Conser
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Tang
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sandrasagra A,
, Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    bronchoconstriction; allergy; hyposecretion; pain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.9%;
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Pred. No. 1
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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nuclei grids associated with lung airway or lung dysfunction, and

bronchodilating

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English

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CC analgesic, hypotensive, immunosuppressive and cyrostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or CC treating a respiratory, lung or malignant disease. The administered CC composition comprises oligo and is administered to reduce the production CC or availability, or to increase the degradation of the target mRNA or to CC reduce the amount of target polypeptide present in the lungs. The CC pulmonary obstruction, and/or bronchoconstriction and/or lung CC inflammation, allergies and/or surfactant hypoproduction are associated CC with a disease or condition such as pulmonary vasoconstriction, respiratory CC distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary CC transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to CC thymidines present in the target RNA serves to prevent the breakdown of CC thymidines present in the target RNA serves to prevent the breakdown of CC thymidines present in the target RNA serves to prevent the breakdown of CC prevent any unwanted effects due to it
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Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA.
                                                                                 Nyce JW,
Miller S,
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                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                   pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PDE4C-derived oligonucleotide SEQ ID 14313.
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                                                                                                                                                                                           24-APR-2001; 2001US-0286036P
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                                                                                   Sandrasagra A,
L, Shahabuddin
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Pred. No. 1.
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); Mismatches
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                                                                                                        Katz
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cc oligonucleotides are derived from a gene encoding or regulating conjuncted these are derived from a gene encoding or regulating conjuncted the target polypeptide associated with lung airway or lung conjuncted in the contrast of the corresponding mRNA. Cc The invention also describes a kit, that comprises: (a) a delivery construction or cancer and can be anti-sense to the corresponding mRNA. Cc The invention also describes a kit, that comprises: (a) a delivery constructions for adding a carrier and for use of the kit. The composition construction for adding a carrier and for use of the kit. The composition conformation construction is useful for preventing or creating a respiratory, lung or malignant disease. The administered composition composition composition is useful for preventing or conformation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The conformary obstruction, and/or bronchoconstriction and/or lung conformary obstruction, and/or bronchoconstriction are associated with a disease or condition such as pulmonary vasoconstriction, are condition such as pulmonary vasoconstriction, conformation, allergies and/or surfactant hypoproduction are associated conformation, allergies, asthma, impeded respiration, respiratory conformation, managed content of the anti-sense oligos corresponding to thymations present in the target RNA serves to prevent the breakdown of the conformation into the system of the sortes of the sortes of the system of the s
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the oligonucleotides into products that free adenosine into e.g., lung, brain, heart, kidney, etc, tissue environment ar prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 15; SEQ ID NO 14313; 763pp; English
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                                                                                                                                                      the system
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Sequence 20 BP; N A; 5 C; 7 <u>ი</u> 0 7, 0 U; 0 Other;

thereby,

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Ś Matches Query Match Local 199 Similarity ATGTTGGTCAGGCTGGTCTC 218 Conservative 1.9%; 0 Pred. Score 18.4; 20 Mismatches No. 1.3e+03; DB 1; Length 20; Indels 0; Gaps 0

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RESULT 815 ADH70951/c 

ADH70951 standard; DNA; 20 ВP

ADH70951;

25-MAR-2004 (first entry)

Human Vbeta PCR primer #95.

rsus host disease; neoplastic disease;

hypersensitivity;

human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease Addison's disease; atrophic gastritts; degenerative nervous system disease; multiple sclerosis; Alzheimer's disease; hypersensitivity; Goodpasture's syndrome; allergy; type II hypersensitivity; Goodpasture's syndrome; allergy; type II hypersensitivity; foodpasture's syndrome; type IV hypersensitivity; leprosy; infectious disease; viral infection; Candida; parasitic infection; schistcosome, filaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain of the control schistosome; viral infection;

Homo sapiens

Pharmaceutical composition for treating asthma, has antisense

WPI; 2003-093058/08

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RESULT 816
ADH54084/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a kit for diagnosing and treating T-cell CC associated diseases which comprises a panel of nucleic acid primers gpecifically priming and allowing amplification of each Vbeta gene, CC VbetaRNA or CDNA. The kit is useful for diagnosing organ transplant C rejection and diagnosing and treating T-cell associated diseases (including autoimmune diseases, degenerative nervous system diseases, CC graft versus host disease, hypersensitivity diseases, infectious disease, CC and neoplastic diseases. Autoimmune diseases include Addison's disease, CC atrophic gastritis. Degenerative nervous system diseases include multiple CC alropsis and Alzheimer's disease. Hypersensitivity diseases include Type CC I hypersensitivities such as those present in CC Goodpasture's syndrome and Type IV hypersensitivities such as those CC caused by viruses such as HIV, fungal infections such as those caused by chast genus Candida, parasitic infections such as those caused by Schistosomes, filaria and bacterial infections such as those caused by wycobacterium. Neoplastic diseases include lymphoproliferative diseases
                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, nucleic acid primers specifically priming and allowing amplific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US2002150891-A1.
              Homo
                                                           human; neurodegenerative disease; urokinase plasminogen activator; uPA; gamma-synuclein; SNCG; insulin degrading enzyme; IDE, kinsein-like protein 1; KNSLI; lySosomal acid lipase; LIPA; tumour necrosis factor receptor SF6; TNFRSF6; Alzheimer's disease; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hood LE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-SEP-1994;
19-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-MAR-1999;
                                                                                                                                      Human neurodegenerative disease-related sequencing primer SeqID211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vbeta gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-059052/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HOOD/)
(ROWE/)
                                            primer; ss; sequencing.
                                                                                                                                                                       25-MAR-2004
                                                                                                                                                                                                                                    ADH54084 standard;
              sapiens
                                                                                                                                                                                                                                                                                                                                               386 CCCAAAGTGCTGGGATTACA 405
                                                                                                                                                                                                                                                                                                                   80
                                                                                                                                                                                                                                                                                                                                                                                19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ROWEN L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          leukaemias,
The present
                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                   CCCAAAGTGCTGGGATTATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rowen L;
                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 5 A; 5 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 1145; 164pp;
                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                       (first entry)
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95US-00531241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9905-00263959
                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                             1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lymphomas and cancers such as cancer of sequence represents a Vbeta PCR primer.
                                                                                                                                                                                                                                    20
                                                                                                                                                                                                                                    ВB
                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                             Score 18.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            English.
                                                                                                                                                                                                                                                                                                                                                                                             1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
                                                                                                                                                                                                                                                                                                                                                                                Indels
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amplification
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ation of a
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RESULT 817
ADI25029
ID ADI250
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                        This invention relates to a novel method of determining a predisposition for or the occurrence of neurodegenerative disease comprising detecting in a target nucleic acid obtained from the subject the presence of an allelic variant of polymorphic regions of human genes selected from urokinase plasminogen activator (upA), gamma-symuclein (SNCG), insulin degrading enzyme (IDE), kinesin-like protein 1 (KNSL1), lysosomal acid lipase (LIPA) and tumour necrosis factor receptor SF6 (TNFRSF6). The method is useful in determining the presence or predisposition to a neurodegenerative disease, particularly Alzheimer's disease. The present sequence is that of a sequencing primer which was used for sequencing of a region of the human KNSL1 gene in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-OCT-2001;
25-OCT-2001;
02-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disease,
presence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Becker KD,
Bertram L,
                                                                        Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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08-NOV-2001;
                                                                                                                                                     Human ZNF9
                                                                                                                                                                                                                                  ADI25029 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Determining a predisposition for or the occurrence of neurodegenerative disease, particularly Alzheimer's disease, comprises determining the presence of a polymorphism in the uPA, SNCG, IDE, KNSL1, LIPA or TNFRSF6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-OCT-2002;
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                                                 WO2004005541-A1
                                                                                                                dominant negative mutant RAB7; dominant negative mutant peripheral neuropathy; human; ZNF9; PCR; primer; ss.
                                                                                                                                                                               22-APR-2004
                        15-JAN-2004.
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                                                                                                                                                                                                                                                                                                                    1064 CGCTAATTTTTGTATTTTCA 1083
                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                                                                    19;
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                                                                                                                                                                                                                                                                                                                                                                                                       20
                                                                                                                                                                                                                                                                                                  CGCTAATTTTTGTATTTTTA 1
                                                                                                                                                     exon 1 forward PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 211; 205pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                      BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Velicelebi G,
Saunders AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001US-0348065P.
2001US-0336983P.
2001US-0336929P.
2001US-0338010P.
2001US-0338363P.
2001US-0338363P.
                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                      12
                                                                                                                                                                                                                                                                                                                                                                                                      A; 2 C;
                                                                                                                                                                                                                                                                                                                                                               1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ellliott KJ, Wang
Mullin KM, Sampson
                                                                                                                                                                                                                                                                                                                                                                                                      N
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                                                                                                                                                                                                                                                                                                                                                   Score 18.4; D
Pred. No. 1.3e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                      <u>و</u>;
                                                                                                                                                                                                                                                                                                                                                                                                      4 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                      0 Other;
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                                                                                                                                                                                                                                                                                                                                                                            Length
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                                                                                                                                                                                                                                                                                                                                                                               20;
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08-JUL-2003; 2003WO-EP050290

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ADH76733/c
ID ADH767
XX
AC ADH767
XX
DT 22-APR
XX
MCHR1
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melani
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obesit
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volesit
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VO2003
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IN-DEC
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VO2003
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PN W02003
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PN W02003
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VO2003
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PN W0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes an isolated nucleic acid (I) coding for a dominant negative, mutant RAB7 polypeptide and/or a dominant negative, mutant RAB7 polypeptide and/or a dominant negative, mutant ARHGEF10 polypeptide. (I) contains in comparison to the wild type RAB7 encoding sequence comprising 624 bp (8B0 ID NO: 1, AD125025) and/or to the wild type ARHGEF10 encoding sequence comprising 3366 bp (SEQ ID NO: 3, AD125027), one or more mutations, where the presence of the nucleic acids is indicative for a predisposition or presence of a peripheral neuropathy. Also described: (1) a nucleic acid probe which is a fragment of (I); (2) a recombinant vector comprising (I); (3) a host cell comprising a recombinant vector of (2); (4) a method for the preparation of a diagnostic assay to detect the presence of a peripheral neuropathy in a human; and (5) a transgenic non-human animal comprising the vector of (2); (I) is useful for isolating and detecting human is useful for detecting the presence of peripheral neuropathy in a human. The diagnostic assay is useful for detecting the presence of peripheral neuropathy in a human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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08-APR-2003; 2003EP-00076033.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                Platzer M,
Reichwald
                                                                                                                                                                                                                                                                                         18-DEC-2003
                                                                                                                                                                                                                                                                                                                                               WO2003104489-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                obesity; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        melanin-concentrating hormone receptor 1; MCHR1; anorectic; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MCHR1 genomic sequence analysis primer #42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            used in an
                                                                                                                                                                                                                                05-JUN-2003; 2003WO-EP005917.
                                                                                                                                                                             05-JUN-2002; 2002EP-00012569.
                                                                                                                    (UYPH-) UNIV PHILIPPS MARBURG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         869 GATTACAGGCGTGAGCCACC 888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           plated nucleic acid coding for a dominant negative, mutant RAB7 stide and/or a dominant negative, mutant ARHGEF10 polypeptide, for detecting the presence of peripheral neuropathy in a human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
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                                   χ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                           Platzer C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INTERUNIVERSITAIR INST BIOTECHNOG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
                                                              Gudermann
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                                                           'n
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                                                                 Hebebrand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Verhoeven
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                 Hinney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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in the specification and at least 8 bases of surrounding sequence of the MCHR1 gene. The composition has anorectic activity. The polynucleotide composition may be used in gene therapy to treat the disorders of the invention. The composition is useful for diagnosing obesity related to the presence of a molecular variant of the MCHR1 gene or a susceptibility to the disorder. The MCHR1 protein or polynucleotide is useful for preparing a medicament for treating or preventing obesity related to the presence of a molecular variant of the MCHR1 gene. This polynucleotide represents an MCHR1 primer of the invention
                                                                                                                                                                                                                                                                    The invention relates to a novel diagnostic polynucleotide composition. The polynucleotide composition comprises: a sequence encoding a polypeptide with defined sequences given in the specification; a sequence capable of hybridizing to a melanin-concentrating hormone receptor 1 (MCHR1) gene; a polynucleotide encoding an MCHR1 polypeptide; or a sequence comprising one or more of the nucleotide exchanges (SNP's) given
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New diagnostic composition, useful for diagnosing obesity related to the presence of a molecular variant of the MCHR1 gene or a susceptibility to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-062377/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Page 43; 76pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             English.
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Query Match
Best Local 9
                                       Matches
                                                                            Sequence 20 BP; 4 A; 3 C; 8 G; 5 T; 0 U; 0 Other;
           541 CCTCAGCCTCCCAAGTAGCT 560
20
                                       19;
                                                Similarity
CCTCAGACTCCCAAGTAGCT 1
                                      Conservative
                                                1.9%;
95.0%;
                                       <u>,,</u>
                                                Score
Pred.
                                        Mismatches
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                                                         18.4;
                                                .3e+03;
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ADH76678
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AX
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single nucleotide polymorphism; anorectic; gene therapy; obesity; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MCHR1 locus
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WO2003104489-A2

18-DEC-2003

05-JUN-2003; 2003WO-EP005917

05-JUN-2002; 2002EP-00012569.

(UYPH-) UNIV PHILIPPS MARBURG

Platzer M, Reichwald Platzer C, Gudermann T, Hebebrand ç Hinney

WPI; 2004-062377/06.

New diagnostic composition, usef presence of a molecular variant the disorder. , useful riant of for the diagnosing obesity related to the or a susceptibility to

Example 1; Page 28; 76pp; English

The invention relates to a novel diagnostic polynucleotide composition.

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Query Match
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The polynucleotide composition comprises: a sequence encoding a sequence polypeptide with defined sequences given in the specification; a sequence capable of hybridizing to a melanin-concentrating hormone receptor 1 (MCHR1) gene; a polynucleotide encoding an MCHR1 polypeptide; or a sequence comprising one or more of the nucleotide exchanges (SNP's) given in the specification and at least 8 bases of surrounding sequence of the MCHR1 gene. The composition has anorectic activity. The polynucleotide composition may be used in gene therapy to treat the disorders of the invention. The composition is useful for diagnosing obesity related to
                                                                                                                                                                                                                                                                                                                                                                                           presence of a molecular the disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                   New diagnostic composition, useful presence of a molecular variant of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Platzer M,
Reichwald
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 single nucleotide polymorphism; anorectic; gene therapy; obesity; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               melanin-concentrating hormone receptor 1; MCHR1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MCHR1 locus
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                                                                                                                                                                                                                                                                                                                               Example 2; Page 45; 76pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYPH-) UNIV PHILIPPS MARBURG
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                                                                                                                                                                                                                                                                  invention relates to a novel diagnostic polynucleotide composition.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Platzer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP primer #41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gudermann
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No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                diagnosing MCHR1 gene
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                                                                                                                                                                                                                                                                                                                                                                                                                   obesity related to the or a susceptibility to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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Similarity

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.3e+03;

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RESULT 821
ADJ46656/c
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Best Local S
Matches 19
                                                                                                             encoding requiem which specifically hybridises with the nucleic acid molecule encoding requiem and inhibits the expression of requiem. The compound, particularly the antisense oligonucleotide is useful in modulating the function of nucleic acid molecules encoding requiem. The antisense compound can also be used as research tools and diagnostics. I can also be used as tools in differential and/or combinatorial analyses to elucidate expression patterns of a portion or the entire complement of genes expressed within cells and tissues. The compound can also be used genes expressed within cells and tissues. The compound can also be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the presence of a molecular variant of the MCHR1 gene or a susceptibility to the disorder. The MCHR1 protein or polymuclectide is useful for preparing a medicament for treating or preventing obesity related to the presence of a molecular variant of the MCHR1 gene. This polymuclectide represents an MCHR1 SNP primer of the invention.
                                               for treating diseases or conditions associated with requiem, preferably hyperproliferative disorder, e.g. cancer or a developmental disorder. compound can also be used as prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence
                                                                                                                                                                                                                                                                                                                     New antisense compound targeted to a nucleic acid molecule encoding requiem, useful for modulating expression of requiem or for treating cancer or developmental disorders.
                                                                                                                                                                                                                                                                                       Example 15; SEQ ID NO 131; 66pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-AUG-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      developmental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human; requiem; hyperproliferative disorder; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human requiem target sequence ISIS #122508.
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                                                                                                                                                                                                                                                       The invention relates to a compound targeted to a nucleic acid molecule
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                                                                                                                                                                                                                                                                                                                                                                                                                            Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS
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                               the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                            Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disorder; infection; inflammation; tumour formation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.9%;
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                               requiem target sequence.
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Pred. No. 1
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                                                                                                   preferably
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RESULT 822
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                                                                              RESULT 823
ADJ59878
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a compound targeted to a nucleic acid molecule encoding requiem which specifically hybridises with the nucleic acid molecule encoding requiem and inhibits the expression of requiem. The compound, particularly the antisense oligonuclectide is useful in modulating the function of nucleic acid molecules encoding requiem. The antisense compound can also be used as research tools and diagnostics. It can also be used as tools in differential and/or combinatorial analyses to elucidate expression patterns of a portion or the entire complement of genes expressed within cells and tissues. The compound can also be used for treating diseases or conditions associated with requiem, preferably hyperproliferative disorder, e.g. cancer or a developmental disorder. The compound can also be used as prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADJ46607 standard;
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-AUG-2002; 2002US-00212993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer or developmental disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bennett CF,
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                                                    ADJ59878 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 15; SEQ ID NO 82; 66pp; English.
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                                                                                                                                                                                                                                                                                                                              Local Similarity
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                                                                                                                                                                                                                           AGCCTCCCAAGTAGCTGGGA 564
                                                                                                                                                                                                                                                                                                                                                                                                                20
                                                                                                                                                                                             AGCCTCTCAAGTAGCTGGGA 20
                                                                                                                                                                                                                                                                                                                                                                                                                BP;
                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Freier SM,
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                                                                                                                                                                                                                                                                                                                                                                                                                5 A;
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                                                                                                                                                                                                                                                                                                                              1.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       requiem
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                                                       20 BP
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                                                                                                                                                                                                                                                                                                                              Score 18.4;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disorder; cancer;
                                                                                                                                                                                                                                                                                                                              1.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                        DB 1; Length 20;
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ADJ59878

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                                                                                                                               RESULT 824
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                                                                                                                                                                                                                                                                                                                                   The present invention relates to an oligonucleotide anti-sense to e.g., cc initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-cc enterleukin (II)-4 receptor, II-5 receptor or salts of the coligonucleotide and optionally surfactant operatively linked to the coligonucleotide. The method is useful for preventing or treating a cc oligonucleotide. The method is useful for preventing or treating a cc respiratory or lung disease, which involves administering to the airways cf a subject an effective amount of an inhibitor. The oligonucleotide is cc of a respiratory or lung disease. The respiratory or lung disease is consent for mairway inflammation, allergy(ies), asthma, impeded cc respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (CODD), allergic rhinitis (AR), acute respiratory distress syndrome cc (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway invention. The present sequence represents an oligonucleotide of the construction.
                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide associated to RANTES #127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2004011613-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; SEQ ID NO 734; 85pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shahabuddin
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            Oligonucleotide associated to
                                                                                                                                                                                                                                                                                               Sequence 20
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                                            06-MAY-2004
                                                                      ADJ59868;
                                                                                                    ADJ59868 standard; DNA;
                                                                                                                                                                                                       641 CACCCAGGCTGGAGTGCAGT 660
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din S, Lu
                                                                                                                                                                            cecccaecrecaerecaer 20
                                                                                                                                                                                                                                                                                               BP;
                                                                                                                                                                                                                                       Conservative
                                            (first entry)
                                                                                                                                                                                                                                                                                               3 A;
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                                                                                                                                                                                                                                                   1.9%;
                                                                                                                                                                                                                                                                                                 6 C; 8
                                                                                                      20
                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                 G; 3 T;
                                                                                                                                                                                                                                       Score 18.4; DB 1
Pred. No. 1.3e+03
0; Mismatches
               RANTES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       receptor;
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                                                                                                                                                                                                                                                                                                  0 U; 0 Other;
                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       lung disease;
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RESULT 825
ADJ59877
ID ADJ598
XX
XX
AC ADJ598
XX
DT 06-MAY
XX
DT 01igon
XX
XX
interl
KW interl
KW airway
XW cystic
KW pulmon
KW ss.
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                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 oligonuclectide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonuclectide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CP), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
             airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                Oligonucleotide associated
                                                                                                                                   06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      oligonucleotide and optionally surfactant operatively linked to oligonucleotide. The method is useful for preventing or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant gene CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-203534/19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-JUL-2002; 2002US-0399076P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          airway inflammation; allergy; asthma; impeded respiration;
cystic fibrosis; acute respiratory distress syndrome;
pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                              interleukin; IL-4 receptor; IL-5 receptor; lung disease;
                                                                                                                                                                  ADJ59877;
                                                                                                                                                                                                   ADJ59877 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            interleukin; IL-4 receptor; IL-5 receptor; lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        initiation codon,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (EPIG-)
                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                     642
                                                                                                                                                                                                                                                                                                                                                      19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EPIGENESIS
                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                       ACCCAGGCTGGAGTGCAGTG 661
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 724;
                                                                                                                                                                                                                                                                                   ACCCAGGCTGAGTGAAGTG
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                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 5 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention relates to an oligonucleotide anti-sense to codon, coding region with 2-10 nucleotides of 5'-end
                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                 (first
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                                                                                                                                                                                                 DNA;
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H, Cong H;
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                                                                                                                                 entry)
                                                                                                                                                                                                                                                                                                                                                                    95.0%;
                                                                                                                                                                                                                                                                                                                                                                                     1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        85pp; English.
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                                                                                                to RANTES
                                                                                                                                                                                                   ВP
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                                                                                                                                                                                                                                                                                                                                                                       Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                   Score 18.4; DB 1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                  #126
                                                                                                                                                                                                                                                                                                                                                                      1.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Miller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease;
                                                                                                                                                                                                                                                                                                                                                      Indels
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RESULT 826
ADJ60947
ID ADJ609
XX ADJ609
XX ADJ609
XX Oligon
DE Oligon
XX interl
KW airway
KW cystic
KW pulmon
KW pulmon
KW pulmon
KW pulmon
KW pulmon
KW pulmon
KW Ss.
XX
PN WO2004
XX
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  end of nucleic acid target comprising gene(8) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonuclectide and optionally surfactant operatively linked to the oligonuclectide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonuclectide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                  interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to an oligonucleotide anti-sense tinitiation codon, coding region with 2-10 nucleotides of 5'-end end of nucleic acid target comprising gene(s) chosen from e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                       Oligonucleotide
                                                                                                                                                                                                                                                                                      06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                   Homo sapiens
                                                                                                                             pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                         ADJ60947
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 BP; 2
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in S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                            associated to PDE4C #13.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;
                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C; 7
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Pred. No. 1.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-FEB-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nyce JW, Tang
Shahabuddin S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-JUL-2002; 2002US-0399076P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-JUL-2003; 2003WO-US023509
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                                                                                                                                                                                                                                                                                                                     airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide associated to RANTES #17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADJ59768 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim
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                                                                                                                                                                                                                                                                                                                                                                                                            interleukin; IL-4 receptor; IL-5 receptor; lung disease;
                                                                                                                                                                                                                                                                                                                  pulmonary hypertension; lung
      29-JUL-2002;
                                                               25-JUL-2003;
                                                                                                                        05-FEB-2004
                                                                                                                                                                              WO2004011613-A2
                                                                                                                                                                                                                                    Homo sapiens.
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      2002US-0399076P
                                                            2003WO-US023509
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Lu H, Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 18.4; DB 1;
Pred. No. 1.3e+03;
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Mismatches

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873 ACAGGCGTGAGCCACCACGC 892

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Query Match
Best Local S
Matches 19
                                                                                                                                       end of nucleic acid target compliancy years.

Interleukin (II)-4 receptor, II-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway inflammation, allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway inflammation, are an airway inflammation.
                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to an oligonucleotide anti-sense to e-
initiation codon, coding region with 2-10 nucleotides of 5'-end and
end of nucleic acid target comprising gene(8) chosen from e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel single or multiple target oligonuclectide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nyce JW, Tang
Shahabuddin S,
                                                                           Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disease e.g., asthma.
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                      Similarity
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                                                                                BP; 4 A;
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Lu
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                    1.9%;
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                                                                                C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cong H;
                                                                                G; 1 T; 0 U; 0 Other;
                        Score 18.4;
Pred. No. 1
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                        .3e+03
                                           DB 1;
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                                           Length
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                                                                                                                                                                                                                                                                     RESULT 828
                                                                                                                                              airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
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 Nyce JW, Tang
Shahabuddin S,
                                             29-JUL-2002; 2002US-0399076P
                                                                                                   WO2004011613-A2
                                                                                                                     Homo
                                                                                                                                        88
                                                                                                                                                                            interleukin; IL-4 receptor; IL-5 receptor;
                                                                                                                                                                                            Oligonucleotide associated to PDE4C
                                                                                                                                                                                                                                  ADJ60990
                                                                                                                                                                                                                                                    ADJ60990
                            (EPIG-)
                                                                 25-JUL-2003; 2003WO-US023509
                                                                                   05-FEB-2004.
                                                                                                                     sapiens
                            EPIGENESIS
                                                                                                                                                                                                                                                    standard; DNA;
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          Tang
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£,
  Sandrasagra A,
H, Cong H;
                              PHARM INC.
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            Aguilar
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                                                                                                                                                                            lung disease;
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RESULT 829
ADJ59767
ID ADJ597
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Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 2; SEQ ID NO 1846; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-203534/19.
 Novel single or multiple target oligonucleotide anti-sense to initiation codons and introns of respiratory disease-relevant
                                                                                                                                                                                                                                                                                                                interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 initiation codon, coding region with 2-10 nucleotides of 5'-envend of nucleic acid target comprising gene(s) chosen from e.g.
                                                                          Nyce JW, Tang
Shahabuddin S,
                                                                                                                                                                                                                                                                                                                                                                                          Oligonucleotide
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                                              WPI; 2004-203534/19
                                                                                                                                                  29-JUL-2002; 2002US-0399076P
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                                                                                                                                                                                                                                          WO2004011613-A2
                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADJ59767 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention.
                                                                                                                      (EPIG-) EPIGENESIS PHARM INC
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                                                                   y L,
Lu
                                                                                                                                                                                                                                                                                                                                                                                          associated to RANTES #16.
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                                                                          Sandrasagra H, Cong H;
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No. 1.3e+03;
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                                                                                      Aguilar
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.
                                                                                           Miller
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e.g.
genes e.g.,
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disease e.g., asthma.
                                                                                                                                            CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
  initiation codon,
                       present
                                                                       2
invention relates to an oligonucleotide anti-sense to e. odon, coding region with 2-10 nucleotides of 5'-end and
                                                                          ij
                                                                          NO 623;
                                                                       85pp; English.
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end of nucleic acid target comprising gene(s) chosen from e.g. ent of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways respiratory or lung disease, which involves administering to the airways respiratory or lung disease, which involves administering to the airways respiratory or lung disease, which involves administering to the airways respiratory or lung disease, which involves administering to the airways respiratory or lung disease, which involves administering to the airways respiratory or lung disease, which involves administering to the airways respiratory or lung disease, which involves administering to the airways respiratory or lung disease, which involves administering to the airways respiratory or lung disease, which involves administering to the airways respiratory or lung disease, which involves administering to the airways respiratory or lung disease, which involves administering to the airways respiratory or lung disease. of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway (ARDS) pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide , airway de of the ωĠ

Sequence 20 BP; 4 ð 5 C; 8 G; 3 T; 0 U; 0 Other;

Matches Query Match Local 1 Similarity
19; Conserv Conservative 1.9%; 0, Score 18.4; DB 1 Pred. No. 1.3e+03 Mismatches DB 1; Length 20; Indels 0 Gaps 0

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868 GGATTACAGGCGTGAGCCAC 887

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RESULT 830

RESULT 830

ADJ59830

ID ADJ598

XX ADJ598

XX ADJ598

XX O1:gon

XX inter1

KW airway

KW cystic

KW pulmor

KW wpi;

KW (EPIG-

KW Nyce |

PI Shahal

KW wpi;

KW pulmor

KW intelled

KW pulmor

KW intelled

KW intelled

KW intelled

KW intelled

KW intelled

KW interled

KW inte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotide associated to RANTES #79.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IL-4 receptor; IL-5 receptor;
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Claim

2

SEQ

ID NO 686;

85pp; English

disease e.g., CCR1, RANTES,

asthma.

nitiation

single or multiple target oligonucleotide anti-sense to ation codons and introns of respiratory disease-relevant RANTES, MCP4, useful for prophylaxis or treating respira

or treating respiratory

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genes

e.g.,

Nyce JW,

(EPIG-)

EPIGENESIS

Shahabuddin

Tang din S,

Lu Lu

Sandrasagra H, Cong H; PHARM INC.

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Aguilar

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Miller

WPI; 2004-203534/19.

05-FEB-2004.

29-JUL-2002; 2002US-0399076P 25-JUL-2003; 2003WO-US023509

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ADJ59829
ID ADJ598
Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome;
The present invention relates to an oligonucleotide anti-sense initiation codon, coding region with 2-10 nucleotides of 5'-enc end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide associated
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                                                                                                                                       Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                     Shahabuddin
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19; Conserv
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                                                                                             SEQ
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in S,
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Lu
                                                                                             NO 685;
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                                                                                                                                                                                                                                   Sandrasagra
H, Cong H;
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                                                                                             85pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                      Miller
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                                          of 5'-end and
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initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is

Claim disease

2; SEQ ID NO 1818; 85pp; English.

invention relates to

an oligonucleotide anti-sense

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RESULT 832
RDSG0962
ID ADJG09
XX ADJG09
XX ADJG09
XX ADJG09
XX Office
DT 06-MAY
XX Interl
KW airway
KW cystic
KW bulmon
KW bulmon
KW BS.

XX Homo s
XX Interl
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           interleukin; IL-4 receptor; IL-5 receptor; lung disease;
airway inflammation; allergy; asthma; impeded respiration;
cystic fibrosis; acute respiratory distress syndrome;
pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                                                                    Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                            WPI; 2004-203534/19
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                                                                                                                                                                          e.g., asthma.
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                                                                                                                                                                                                                                                                       Tang L,
din S, Lu
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                                                                                                                                                                                                                                                                       Sandrasagra
H, Cong H;
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No. 1
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RESULT 833
ADJ59773
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Best Local S
Matches 19
               The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (II)-4 receptor, II-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway the treatment of the control of the con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nyce JW, Tang
Shahabuddin S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide associated to RANTES
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                                                                                                                                                                                                                                                                                                                                                                                                                                    disease e.g., asthma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
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The present sequence represents
                                                                                                                                                                                                                                                                                                                                                                                         ID NO 629;
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H, Cong H;
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95.0%;
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                                                                             end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (II)-4 receptor, II-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(les), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases.
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airway inflammation; allergy; asthma; impeded respiration;
cystic fibrosis; acute respiratory distress syndrome;
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Shahabuddin
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Sequence
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                                                                                                                                                                                                                                                                   Claim
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                                      (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airwibstruction. The present sequence represents an oligonucleotide of
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din S, Lu
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Query Match

Score 18.4;

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Pred. No.

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Mismatches

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RESULT 835
ADJ59880
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                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 mucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (II)-4 receptor, II-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is the control of treatment of a subject and the control of treatment of a subject and the control of treatment of a subject and the control of treatment of the control of the control of treatment of the control o
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                                                                                                                                                                                                                     useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CP), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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Shahabuddin
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                                                                                                                                                      Sequence
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RESULT 837 ADJ96297

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RESULT 8366

RESULT 8366

RESULT 8366

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PT C
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CC interleukin (II)-4 receptor, II-5 receptor or salts of the coligonucleotide and optionally surfactant operatively linked to the coligonucleotide. The method is useful for preventing or treating a cC respiratory or lung disease, which involves administering to the airways cf a subject an effective amount of an inhibitor. The oligonucleotide is cc useful for production of a medicament for the prevention and/or treatment cc arepiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded cc (Copp), allergic rhintis (AR), acute respiratory distress syndrome (CD), allergic rhintis (AR), acute respiratory distress syndrome cc (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway construction. The present sequence represents an oligonucleotide of the
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Shahabuddin S,
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                                                                                                                                             Similarity
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                                                            ATGITGGTCAGGCTGGTCTC 218
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      ATGTTGGCCAGGCTGGTCTC 20
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                                                                                                                                                                            Score 18.4;
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Page 424
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RESULT 838
ADJ96333/c
ID ADJ963
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XX ADJ963
XX O6-MAY
XX Human
XX Breast
KW Breast
KW Breast
KW antise
XX Synthe
OS Uniden
XX US2004
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Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                          New antisense oligonucleotide compounds, useful for diagnosing, preventing and/or treating conditions with aberrant activity of breast cancer-1, such as breast, ovary, prostate and/or peritoneum cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense therapy; antisense;
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  US2004014051-A1
                                                                                                                                                                                                                                                               Sequence 20
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                    Synthetic.
Unidentified
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                                                         Breast
                                                                            Human breast cancer-1 associated antisense oligonucleotide #51.
                                                                                               06-MAY-2004
                                                                                                                                     ADJ96333 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC.
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                                                         cancer-1; diagnosis; hyperproliferative disorder; cancer;
                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                    CTCTGTCGCCCAGGCTGGAG 20
                                                therapy;
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                                                                                              (first entry)
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Pred. No. 1
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RESULT 839
ADJ96393
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense oligonucleotide compounds, useful for diagnosing, preventing and/or treating conditions with aberrant activity of breast cancer-1, such as breast, ovary, prostate and/or peritoneum cancers.
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US2004014051-A1.
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                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                         antisense therapy;
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                                                                                                                                                                                                                                                                                           modified_base
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Similarity 95.0%;
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/note= "2'- metho
16. .20
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/note= "Phosphorothioate backbone where all cytidines are
5'- methylcytidines"
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Pred. No. 1.
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RESULT 840
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XX ADJ964
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XX Breast
KW IS2004
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Best Local (
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                            New antisense oligonucleotide compounds, useful for diagnosing, preventing and/or treating conditions with aberrant activity of breast cancer-1, such as breast, ovary, prostate and/or peritoneum cancers.
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Pred. No. 1.
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RESULT 841
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Best Local
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                                                                                                                                                                                                                                                                                                                                                  08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
07-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention is directed to novel antisense compounds targetted to breast cancer-1 proteins and their encoding nucleic acids. The invention is useful for the diagnosis, prevention and/or treatment of diseases and conditions associated with aberrant expression and activity of breast cancer-1 such as a hyperproliferative disorder in particular breast, ovary, prostate and peritoneum cancers. The invention is also used in antisense therapy. The present sequence is human breast cancer-1
                                     This invention relates to a novel primers useful for synthesising full length cDNA molecules that encode human proteins. Specifically, it refers to secretory or membrane proteins that are potential therapeutic agents/ target molecules in the field of medicine, and in particular genes encoding proteins that are associated with signal transduction, glycoproteins and transcription. The present invention describes a method for efficiently cloning a full length human cDNA from both the 5' and 3' ends using the oligo-capping method. This oligonucleotide sequence is a human clone specific PCR primer used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 4 A; 7 C; 7 G; 2 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          target oligonucleotide.
                                                                                                                                                                                                                                                    WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oligo-capping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human; medicine; signal transduction; glycoprotein; transcription;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Clone specific
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                                                                                                                                                                              Example 18; SEQ ID NO 4367; 1340pp; English.
                                                                                                                                                                                                           New oligonucleotide primers length human cDNAs.
                                                                                                                                                                                                                                                                                Wakamatsu
                                                                                                                                                                                                                                                                                              Ota T,
                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUL-2000; 2003EP-00025638
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                                                                                                                                                                                                                                                                                              Nishikawa T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard;
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                                                                                                                                                                                                                                                                                                                                                  ; 99JP-00194486.
; 2000JP-00118774.
; 2000JP-00183865.
; 2000EP-00114089.
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                                                                                                                                                                                                                                                                                Sugiyama
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
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95.0%;
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a T, Nagai
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                                                                                                                                                                                                                          (830 cDNAs)
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Pred. No. 1.3e+03
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                                                                                                                                                                                                                                                                                Hayashi K,
i K, Kojima
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S, Otsuki
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           which normally participates in embryonic development, cell differentiation and stimulates apoptosis via caspase-3 activation. The present invention describes antisense oligonucleotides that comprise a
                                           This invention relates to a novel antisense compounds that modulate the expression of human geranyleranyl diphosphate synthase 1 (also known as GGPS1, geranylegranyl pyrophosphate synthatese, GGPPS, gyppsase and geranyltranstransferase) and located on chromosome 1p43. Specifically, refers to compositions useful for inhibiting the expression of GGPS1, which compositions useful for inhibiting the corposations.
                                                                                                                                                    New antisense oligonucleotides for modulating geranylgeranyl synthase 1 expression, useful for diagnosing, preventing or t conditions associated with the protein, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense; ss; human; geranylgeranyl diphosphate synthase 1; GGPS1; geranylgeranyl pyrophosphate synthetase; GGPS; ggppsase; geranyltranstransferase; embryonic development; cell differentiation; apoptosis; 2' MOE wing; phosphorothioate backbone; developmental;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADJ10322 standard;
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                                                                                                                           Example 15;
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 modified sugar moiety, a 2'-O-methoxyethyl
                                                                                                                                                                                                                                Bennett CF,
                                                                                                                            SEQ
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                                                                                                                              ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                      cytidine nucleobases 16. .20
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/note= "OTHER= 2'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disorder;
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                                                                                                                           71; 76pp; English.
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                    methoxyethyl (2' MOE) nucleotides.
ses are 5-methylcytidine."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                              yethyl (2' MOE) nucleotides. All
5-methylcytidine."
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Best Loc Matches

Local

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Pred. No. 1.3e ); Mismatches Score 18.4;

1.3e+03

DB 1;

Length

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Indels

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Gaps

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95.0%;

1.9%;

Query Match

Sequence

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RESULT 843
ADO17893/c
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                                    The present invention relates to a panel of two or more single nucleotide polymorphisms, where each of the polymorphisms of the panel are selected from single nucleotide polymorphisms that are not genetically linked with respect to one another, and where each of the polymorphisms of the panel are selected from single nucleotide polymorphisms that are located outside tandem repeat nucleic acid sequences. The known sample is from a family member. The compromised nucleic acid sequences comprises nucleic acid fragments from 10-100 nucleotides in length. The identity of the one or more single nucleotide polymorphisms is determined using a single base primer extension reaction. The present sequence represents a primer of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  least one modified nucleobase, a 5-methylcytosine. Accordingly, these compounds are useful for treating a disease or condition associated with GGPS1 such as a developmental or hyperproliferative disorder (e.g. cancer) that arise as a result of aberrant apoptosis. As such, these compositions exhibit cytostatic activity and are useful for diagnostics, prophylaxis, research reagents and various kits. This oligonucleotide sequence is a phosphorothicate antisense DNA oligo used to modulate human geranylgeranyl diphosphate synthase 1 expression in an exemplification of
                                                                                                                                                                                                                                                acid
                                                                                                                                                                                                                                                            New panel of single nucleotide polymorphisms comprising two or more single nucleotide polymorphisms, useful for analyzing compromised nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single nucleotide polymorphism;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the invention.
                                                                                                                                                                                                                     Disclosure; SEQ ID NO 120;
                                                                                                                                                                                                                                                                                                                                                               (ORCH-) ORCHID BIOSCIENCES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
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95.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer; ss.
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RESULT 845
ADM14163/c
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AC ADM141
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XX
DF Human
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ADO18178/c
ID ADO18178 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New panel of single nucleotide polymorphisms comprising two or more single nucleotide polymorphisms, useful for analyzing compromised nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
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Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:350.

Query Match Best Local

Similarity

1.9%; 95.0%; 6 C; 5

Score 18.4; Pred. No. 1

1.3e+03; DB 1;

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The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The Chuman mpGES-1 gene is located on chromosome 9, more specifically to Q34.3. The present invention also describes: (1) antisense compounds, CC 934.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1 in cells or tissues; and (3) a method of tracting the expression of CC mpGES-1 in cells or tissues; and (3) a method of tracting an animal CC having a disease or condition associated with mPGES-1. MPGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antifilammatory, neuroprotective, nootropic, antiarthritic, vasotropic, CC antifilammatory, immunomodulator, cardiant, neuroprotective, vasotropic, CC ophthalmological, immunomodulatory and cardiovascular activities, and can CC be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of
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Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                 Gierse JK
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                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chimeric; antisense oligonucleotide; phosphorothioate; human;
microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human mPGES-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADM14236;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ophthalmic,
                                                                                                      25-SEP-2003;
                                                                                                                                                                 WO2004028458-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                        25-SEP-2002;
                                            (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  727
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immunological,
                                                                          2002US-0413549P
                                                                                                      2003WO-US030374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chimeric antisense oligonucleotide SEQ ID NO:423
                                                                                                                                                                                                                                         /note= "
                                                                                                                                                                                                                                                                                                                  residues are
                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                       _mod_
                                                                                                                                                                                                                          /*tag=
                                                                                                                                                                                                                                                                                     /*tag= a
                                                                                                                                                                                                                                                                                                                 note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.9%;
                                                                                                                                                                                           base= OTHER
== "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                       _base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; 2
                                                                                                                                                                                                                                                      "2'-0-methocyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>,,</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 18.4;
Pred. No. 1.
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mPGES-1

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sequence represents a chimeric antisense oligonucleotide human microsomal prostaglandin E2 synthase (mPGES-1). Th

gene is located on chromosome 9, more specifically to resent invention also describes: (1) antisense compounds,

The present invention also

Claim

4; SEQ

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NO 423; 132pp;

English

encoding mPGES-1, useful f inflammation, Alzheimer's

useful

having a sequence targeted to a nucleic acid l for preparing a composition for treating e.g. disease, arthritis, diabetes, cancer or

e.g.,

New antisense compound,

CC antisense or condition associated with mpGES-1 and can be used as mpGES-1, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibits; its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, antiarthritic, vasotropic, CC ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.

Query Match Best Local

Sequence 20 BP; 3 A;

6 C; 5 G; 6 T; 0 U; 0 Other;

Matches

Local Similarity nes 19; Conserv

Conservative

<u>,</u>

Mismatches

1.9%;

Score 18.4; Pred. No. 1.

1.3e+03 DB 1;

Length 20

0,

Gaps

0

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RESULT 848
ADM14395/c
ID ADM143
XX
AC ADM147
XX
XX
DT 01-UU
DT 01-UU
XX
DB Human
XX
DB Human
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U chime
KW chime
KW micro
KW micro
KW immur
KW immu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:582
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                 modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disorder;
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                                                                                                                                                                                                                        /note= "phosphorothioate linkages
residues are 5-methylcytidines"
                                                                                                                                                                                                                             residues are
                                                                                                                                                                                                                                                                                                      _pom/
                                                        'mod_base= OTHER
'note= "2'-O-methocyethyls"
                                                                                                                                                                                                                                                                                                      base= OTHER
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neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia;

reperfusion injury;

ophthalmic disorder;

chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory;

chimeric antisense oligonucleotide SEQ ID NO:1550

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CC necessary immunomodulatory and in gene therapy. The antisense compound condition associated with mpGES-1 chimeric condition associated with mpGES-1 chimeric continuation, neuroprotective, nootropic, antisties, and can be used as mpGES-1 inhibitors and antisense compounds have cytostatic, condition associated with mpGES-1 chimeric contininflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition associated with mpGES-1 e.g., inflammation injury, or condition injury, or 
                                                                                                                                                                                                                                                                                                                       RESULT 849
ADM15363/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9034.3. The present invention in the present invention in the present invention in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense compound, having a sequence targeted to a nucleic acencoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunological,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IJ
                                                                                                                                                                                                                   (first entry)
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== "2'-O-methoxyethyls"
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RESULT 850
ADM15261/c
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                                                                                                                                                                                                         The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 Synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, inhibits its expression; (2) a method of inhibiting the expression; (2) a method of inhibiting the expression of mpGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic, and cantisense oligonucleotides and antisense compounds have cytostatic, ophthalmological, immunomodulator, cardiant, neuroprotective, antisense compound can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                       Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense compound, having a sequence targeted to a nucleic avencoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4;
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                                                                                                                                     Similarity
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                                                               CTCAGCCTCCTGAGTAGCTG 1
                                                                                          CCCAGCCTCCTGAGTAGCTG 736
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                                                                                                                        Conservative
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1. .20
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residues are 5-methylcytidines"
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/mod_ba
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/mod_base= OTHER
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== "2'-O-methocyethyls"
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Pred. No. 1.3e
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                                                                                         The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective.
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antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's
                                                                                                                                                                                                                                                                                                                     Claim 4; SEQ ID NO 1448; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic acid encoding mPGBS-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-305094/28
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/note= "2'-O-methocyethyls"
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/note= "phosphorothioate_linkages and all cytidine
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disease, arthritis, ophthalmic, immunol

hritis, diabetes, immunological, car

es, cancer, ischaemia or reperfusion injury, cardiovascular or neurological disorder.

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Sequence 20

BP; 4 A; 5 C; 8

G; 3

T; 0 U; 0 Other;

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RESULT 851
ADM15038/c
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Best Local Sim
Matches 19;
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                                                                                                           New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                             Claim
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                                NO 1225; 132pp; English
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95.0%;
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e= "2'-O-methoxyethyls"
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Pred. No. 1.
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RRSULT 852
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 20;
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25-SEP-2003; 2003WO-US030374

08-APR-2004.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 19
                                                                                                                                                                                                             chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; SEQ ID NO 1370; 132pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-305094/28.
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                                                                                                                                                    Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                       Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1391.
                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADM15204 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CCTGAGTAGCTGGGATTACA 1
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                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                        1. .20
                  /*tag=
                                                        Location/Qualifiers
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; 6 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 20;
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SXEXEXE

01-JUL-2004 ADM15266;

(first entry)

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chimeric; antisense oligonucleotide; phosphorothioate; human; Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1453

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RESULT 854
ADM15266/c
ID ADM152
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                                                                                                                                                                                                                                     The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The CC human mPGES-1 gene is located on chromosome 9, more specifically to CC 9d34.3. The present invention also describes: (1) antisense compounds, expression; (2) a method of inhibits expression; (2) a method of inhibiting the expression; (2) a method of inhibiting the expression of CC mPGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mPGES-1. MPGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, antiense oligonucleotides and antisense compounds have cytostatic, contininflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, antiinflammatory, neuroprotective, nootropic, antiarthrities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's CC ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                    Matches
                                                                                                                                                                                   Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gierse JK;
                                                                                                                                                                                                                                   Sequence 20 BP; 2 A; 5 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-305094/28
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                  ADM15266
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                                                                                                                                  843 CCTGCCTCGGCCTCCCAAAG 862
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                                                                                                                                                                    19;
                                                                                                                                                                                     Similarity
                  standard;
                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NO 1391; 132pp; English.
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/note= "2'-O-methocyethyls"
i6. .20
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residues are 5-methylcytidines"
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                                                                                                                                                                                   1.9%;
                  DNA;
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e= "2'-O-methoxyethyls"
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                                                                                                                                                                                   Score 18.4; DB 1; Length 20; Pred. No. 1.3e+03;
                                                                                                                                                                       Mismatches
                                                                                                                                                                      Indels
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                                                                                                                                                                    Gaps
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The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to CC 9934.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC mpGES-1 in cells or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonuclectides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, antiaritritic, vasotropic, cophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition associated with mpGES-1 condition injury, or condition injury, or condition associated with mpGES-1 condition injury, or condition injury injury.
Query Match
Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; SEQ ID NO 1453; 132pp; English.
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                                                                                                                                            ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sapiens.
     l Similarity
19; Conser
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                                                                                                                                               immunological,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                residues are 5-methylcytidines"
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/mod_base= OTHER
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                                                                                               4 C; 8
                         1.9%;
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                                                                                                                                               cardiovascular or neurological disorder
                                                                                               G; 3 T; 0 U; 0 Other;
     Score 18.4; I
Pred. No. 1.3e
0; Mismatches
                           .3e+03
                                                  DB 1; Length 20;
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Indels

0

Gaps

0;

The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mBGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric

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RESULT 855
ADM13950/c
ID ADM139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1 imPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                               New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.c inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human mPGES-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                               Claim 4;
                                                                                                                                                                                              WPI; 2004-305094/28
                                                                                                                                                                                                                                           (PHAA ) PHARMACIA CORP
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                                                                                                               SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                               ID NO 137; 132pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chimeric antisense oligonucleotide SEQ ID NO:137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disorder; neurological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                 /note=
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                        residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "phosphorothicate linkages and all cytidine
                                                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "2′-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                           'mod_base= OTHER
'note= "2'-O-methocyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER
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ADM14044/c
ID ADM140
XX
AC ADM140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 856
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostraglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostraglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                     WPI; 2004-305094/28
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                                                                                                                                                          (PHAA )
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'note= "2'-O-methocyethyls'
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Pred. No. 1.
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Query Match

Sequence 20 BP; 5 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

DB 1;

Length 20;

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antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antiinflammatory, neuroprotective, nootropic, antiinflammatory, neuroprotective, nootropic, antiinflammatory, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alabeimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                  The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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ADM14120/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               disorder; neurological disorder; ss.
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/*tag=
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residues are 5-methylcytidines"
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Pred. No. 1.
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base= OTHER

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RESULT 858
ADM14121/c
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                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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         neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy, inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                  chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory;
                                                                                                                                                                                                             ADM14121
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                                                                                                                                   Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:308
                                                                                                                                                                                                                                                                                                    991
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                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                             standard;
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                                                                                                                                                                                                                                                                           CTCCCGGGTTCAAGCGATTC
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95.0%;
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Pred. No. 1.
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RESULT 859 ADM15337/c ID ADM153 XX

ADM15337 standard; DNA;

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                                                                                                                                                          CC sparing a sequence comprising 8-30 bp targeted to a nucleic acid encoding compGES-1, which specifically hybridise with the nucleic acid encoding cc inhibits its expression; (2) a method of inhibiting the expression of cc inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal cc antisense oligonucleotides and antisense compounds have cytostatic, cantidabetic, immunomodulator, cardiant, neuroprotective, and can compitable compounds have cytostatic, cc antidabetic, immunomodulator, cardiant, neuroprotective, asotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can cc ophthalmological, immunomodulator, and cardiovascular activities, and can cc condition associated with mpGES-1 inflammation, Alzheimer's c condition associated with mpGES-1 e.g., inflammation, Alzheimer's componing composition for treating a disease or componing composition injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                           Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human mPGES-1
                                                                                                                                                                                                                                                                                                                                                                                                                        targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; SEQ ID NO 308; 132pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-305094/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-SEP-2002; 2002US-0413549P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        iomo sapiens.
                                                                                                                           Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                           The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (PHAA ) PHARMACIA CORP
 20
                                                             l Similarity
                                                                                                                                                                                                                                                                                                                                                                                          sent sequence represents a chimeric antisense oligonucleotide d to human microsomal prostaglandin E2 synthase (mPGES-1). The PGES-1 gene is located on chromosome 9, more specifically to The present invention also describes: (1) antisense compounds,
                                TCCCGGGCTCAAGCGATTCT 1011
                                                                                                                             BP; 6 A; 5 C; 6 G; 3 T; 0 U;
                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= c
/mod_base= OTHER
/mote= "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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/mod_base= OTHER
/note= "2'-O-methocyethyls"
                                                                               95.0%;
                                                              0,
                                                                                              Score 18.4;
                                                                               Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English
                                                                Mismatches
                                                                               No.
                                                                              .3e+03
                                                                                                                                0 Other;
                                                                                                DB 1;
                                                                                             Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer or
                                                                  <u>,</u>
                                                                  Gaps
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antidiabetic, immunomodulator, cardiant, neuroprotective, assotropic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                        The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9344.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression, (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antissues of condition associated with mPGES-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notiropic; ophthalmological; neuroprotective; notiropic; ophthalmological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADM15337;
                                                                                                                                                                                                                                                                                                                                                     Claim 4; SEQ ID NO 1524; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-305094/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-SEP-2003; 2003WO-US030374
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2004028458-A2
                                                                                                                                            antisense oligonucleotides and antisense compounds have cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mmunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antisense oligonucleotide; phosphorothioate; human;
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es are 5-methylcytidines"
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 cardiovascular
neurological
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RESULT 860
ADM15320/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antinflammatory; neuroprotective; noctropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chimeric; antisense oligonucleotide; phosphorothicate;
                               Claim 4;
                                                             New antisense compound, having a sequence targeted to a nucleic avenceding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                 Gierse
                                                                                                                                                                            25-SEP-2002; 2002US-0413549P
                                                                                                                                                                                                                       08-APR-2004
                                                                                                                                                                                                                                             WO2004028458-A2
                                                                                                                                                                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                                                                                           modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1507
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADM15320;
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                                                                                                           WPI; 2004-305094/28
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                                                                                                                                                     (FHAA)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity
19; Conserv
                                                                                                                                                      PHARMACIA CORP
                              SEQ ID NO 1507; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                  2003WO-US030374.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                        /note= "phosphorothioate linkages residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                mod base= OTHER
                                                                                                                                                                                                                                                                   note=
                                                                                                                                                                                                                                                                                                                                     *tag= a
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Pred. No. 1.3e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                     and
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                                                                                                                                                                                                                                                                                                                                                                      all cytidine
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                                                                          e.g.,
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The present targeted to

sequence represents a chimeric human microsomal prostaglandin

antisense oligonucleotide E2 synthase (mPGES-1). The

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ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.

Sequence 20

BP; 7 A; 7 C; 1 G; 5 T; 0 U;

0 Other; DB 1;

Similarity

1.9%;

Score 18.4; Pred. No. 1.

1.3e+03;

Length 20;

antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotrantiinflammatory, neuroprotective, nootropic, antiarthritic, neuroprotective, nootropic, antiarthritic, neuroprotective, nootropic, antiarthritic, neuroprotective, neuropr

vasotropic ties, and c

human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric

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RESULT 86
RADM13895/
ID ADM11
XX ADM11
XX ADM1
XX ADM1
XX Chim
AC ADM1
XX AC ADM1
XX AC ADM1
XX AC ADM1
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                                                                                                                                                                                                                                                                                                                                                                                                                                           microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; lamunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chimeric; antisense oligonucleotide; phosphorothioate; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADM13895 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                              cardiovascular
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                                                                                                                                                                           modified_base
   25-SEP-2002;
                                  25-SEP-2003;
                                                                  08-APR-2004
                                                                                                 WO2004028458-A2
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   2002US-0413549P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                   2003WO-US030374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chimeric antisense oligonucleotide SEQ ID NO:82
                                                                                                                                                                                                                                                                                                                                                                                                                            disorder; neurological disorder; ss.
                                                                                                                                                                              /note=
                                                                                                                                                                                                                                                             /note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                     'mod_base= Оїньк
'note= "2'-0-methocyethyls"
                                                                                                                                                                                                                            /*tag= a
                                                                                                                                                                                                                                                                                             mod_base= OTHER
                                                                                                                                  note=
                                                                                                                                                                                . 20
                                                                                                                              _base= OTHER
== "2'-O-methoxyethyls'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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The present sequence represents a chimeric antisense oligonucleotide cc targeted to human microsomal prostaglandin E2 synthase (mPCES-1). The CC human mPCES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, cc having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPCES-1, which specifically hybridise with the nucleic acid encoding cc mPCES-1 in cells or tissues; and (3) a method of tracting an animal cc mPCES-1 in cells or tissues; and (3) a method of tracting an animal cc mpcES-1 in cells or tissues; and (3) a method of tracting an animal cc antisense oligonucleotides and antisense compounds have cytostatic, cc antidiabetic, immunomodulator, cardiant, neuroprotective, and can cc ophthalmological, immunomodulatory and cardiovascular activities, and can cc can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's contacting a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's contacting a disease or condition associated with mPGES-1 e.g., inflammation reperfusion injury, or condition associated with mPGES-1 e.g., inflammation, Alzheimer's contacting a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition associated with mPGES-1 e.g., inflamm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound, having a sequence targeted to a nucleic acencoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gierse JK;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ischemia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4; SEQ ID NO 82; 132pp; English.
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e.g.,

닭 S Matches 19; Query Match Sequence 20 BP; 4 A; 4 C; 10 G; 2 T; 0 U; Local 684 CCTCTGCCTCCCGGGTTCAA 703 20 Similarity Conservative 1.9%; 0; Mismatches Score 18.4; DB 1; Pred. No. 1.3e+03; DB 1; 0 Other; Indels 0; Gaps

0

ophthalmic,

immunological, cardiovascular or neurological disorder

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neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                      chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                         01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                          ADM14082 standard; DNA; 20
                                                             modified_base
                                                                                                              Synthetic.
                                                                                                                             Homo
                                                                                                                                                                                                                                                                                                                         Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:269
                                                                                                                           sapiens.
                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                             Location/Qualifiers
                                mod_base= OTHER
                                                                . 20
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/note= "phosphorothioate linkages residues are 5-methylcytidines"

and a11

cytidine

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RESULT 863
ADM14445/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mbGES-1 gene is located on chromosome 9, more specifically to 20 4934.3. The present invention also describes: (1) antisense compounds, C mpGES-1, which specifically hybridise with the nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and C inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal c may be a modern of the expression of mpGES-1 in cells or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, or antininflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or cardiant, in the composition of the
                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 19
microsomal prostaglandin microsomal prostaglandin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
                                             chimeric;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; SEQ ID NO 269; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-305094/28.
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                                                                                       Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:632
                                                                                                                                     01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-SEP-2003; 2003WO-US030374.
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                                                                                                                                                                                                                         ADM14445 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-SEP-2002; 2002US-0413549P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ischemia.
                                                                                                                                                                                                                                                                                                                                                                                 993 CCCGGGCTCAAGCGATTCTC 1012
                                                                                                                                                                                                                                                                                                                                     20
                                                                                                                                                                                                                                                                                                                                                                                                                            19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                     CCCGGGTTCAAGCGATTCTC
                                         antisense oligonucleotide; phosphorothicate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunological, cardiovascular or neurological disorder
                                                                                                                                  (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
/note= "2'-O-methocyethyls"
16..20
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/mod_ba
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                             1.9%;
                                                                                                                                                                                                                         DNA;
                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         base= OTHER
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E 2
                                                                                                                                                                                                                         ВP
synthase; mPGES-1; mPGES-1 inhibitor;
synthase inhibitor; cytostatic; antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                          <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 18.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                   1.3e+03;
1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                        0
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The CC human mPGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, CC mPGES-1, which specifically hybridise with the nucleic acid encoding comprise a sequence comprising 8-30 bp targeted to a nucleic acid encoding compGES-1, which specifically hybridise with the nucleic acid mPGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of compGES-1 in cells or tissues; and (3) a method of treating an animal compGES-1 in cells or condition associated with mPGES-1. MPGES-1 chimeric cantidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, antiarthritic, vasotropic, capthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's composition disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
 Matches
                 Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immunomodulator; cardiant; neuroprotective; antiinflammatory;
neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological;
immunomodulatory; cardiovascular; gene therapy; inflammation;
Alzheimer's disease; arthritis; diabetes; cancer; ischaemia;
reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gierse JK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        reperfusion in cardiovascular
                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-SEP-2002; 2002US-0413549P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                4;
                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ
                                                                   20
                                                                   B₽;
 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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                                                                   P,
                 1.9%;
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== "2'-O-methoxyethyls"
                                                                   7 C; 4 G; 6 T;
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 <u>,,</u>
               Score 18.4; DB 1;
Pred. No. 1.3e+03;
   Mismatches
                                                                   0
U;
                                                                     0 Other;
                                 Length 20;
     Indels
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Gaps
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387

CCAAAGTGCTGGGATTACAG

406

0

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20

CCAAAGTGCTGGGATGACAG 1

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RESULT 864
ADM14651/c
microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; noctropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human mPGES-1 chimeric antisense oligonucleotide SEQ ID
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        The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUL-2004
                                                                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                       25-SEP-2002; 2002US-0413549P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chimeric; antisense oligonucleotide; phosphorothioate;
                                                                                                                                                                                                                                                                                                          WPI; 2004-305094/28
                                                                                                                                                                                            Claim 4; SEQ ID
                                                                                                                                                                                                                                                                                                                                                                        (PHAA ) PHARMACIA CORP.
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16. .20
                                                                                                                                                                                          NO 838; 132pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mod
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      _base= OTHER
e= "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "2'-O-methocyethyls"
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immunomodulator, cardiant, neuroprotective

New antisense compound, having a sequence targeted to a nucleic encoding mPGES-1, useful for preparing a composition for treating the second of the second o

e.g.,

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RESULT 865
ADM15095/c
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; mtcrosomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; halzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human
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                                                             Gierse
                                                                                                                                       25-SEP-2003; 2003WO-US030374
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                                                                                                                                                                08-APR-2004.
                                                                                       (PHAA ) PHARMACIA CORP
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                                      2004-305094/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mPGES-1 chimeric
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunological, cardiovascular or neurological disorder
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                                                                                                                2002US-0413549P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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1. .20
                                                                                                                                                                                                                                                                               /*tag= a
/mod_base=
                                                                                                                                                                                                                                                                                                                      /note= "phosphorothicate linkages residues are 5-methylcytidines"
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16. .20
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                                                                                                                                                                                                                               mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.9%;
                                                                                                                                                                                                                                                                                                                                                  base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antisense oligonucleotide SEQ ID NO:1282
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Pred.
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No. 1
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                                                                                                                                                                                                                                                                                                                                      cytidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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RESULT 866
ADM15230/c
ID ADM152
XX ADM152
XX ADM152
XX OADM152
XX OADM153
XX 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADM15230 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammation,
                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                     modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1417.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       726
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunological,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            g
                                                                                                                                              /note= '
                                                                                                                                                                                                                                                                                                                               residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                           /note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag=
                                                                                                         'tag=
                                                                                                                                                                                                                                                                                                                                                                                                          mod
                                                                                                                                                                                                                     mod_base= OTHER
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                                                                              base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                  base= OTHER
                                     "2'-O-methoxyethyls"
                                                                                                                                                                                    "2'-O-methocyethyls"
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RESULT 867
ADM1471/c
ID ADM144
AC ADM144
AC ADM144
XX
DT 01-JUL
XX
Chimen
DE Human
XX
Chimen
KW micros
KW micros
KW immuno
KW immuno
KW immuno
KW immuno
KW immuno
KW reperi

chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin %2 synthase; mpGES-1 imhibitor; microsomal prostaglandin %2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; immunomodulator; contropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;

Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:658.

01-JUL-2004 ADM14471;

(first entry)

ADM14471 standard; DNA;

20

₽P

Homo sapiens cardiovascular

disorder;

neurological

밁 S

Matches

l Similarity
19; Conserv

Conservative

0

Mismatches

0

Gaps

0

1.3e+03

95.0%;

773 20

TGTATTTTAGTAGAGATGG 792 TGTATTTTAGTAGAGACGG 1

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CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding compGES-1, which specifically hybridise with the nucleic acid encoding compGES-1, which specifically hybridise with the nucleic acid encoding compGES-1 in cells or tissues; and (3) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal compounds and in the mace of condition associated with mpGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, cantidiabetic, immunomodulator, cardiant, neuroprotective, or antidiabetic, immunomodulator, and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound con be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's componition of the disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating exinflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                Sequence 20 BP; 8 A; 6 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                         targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; SEQ ID NO 1417; 132pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-305094/28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence represents a chimeric
                    1.9%;
Score 18.4;
Pred. No. 1.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antisense oligonucleotide
                       Length 20;
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e.g.,

01-JUL-2004 (first entry)

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RESULT 868
ADM15203/c
ID ADM15203 &
XX
AC ADM15203;
XX
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                                                                                                                                                                                   Matches
                                                                                                                                                                                                  Query Match
Best Local
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; SEQ ID NO 658; 132pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-SEP-2003; 2003WO-US030374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-APR-2004
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                                                                                                                                           1060 ACCCCGCTAATTTTTGTATT 1079
                                                                                                                           20
                                                                                                                                                                                   1 Similarity
19; Conserv
                                                   standard; DNA; 20
                                                                                                                                                                                                                                                BP; 9 A; 2 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "phosphorothioate linkages
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mod_base= OTHER
                                                                                                                                                                                             1.9%;
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                                                   BP.
                                                                                                                                                                                   0;
                                                                                                                                                                                                   Score 18.4;
Pred. No. 1.
                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                     1.3e+03
                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                Length
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                                                                                                                                                                                       Indels
                                                                                                                                                                                                                   20;
                                                                                                                                                                                     0
                                                                                                                                                                                     Gaps
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The present sequence represents a chimeric antisense oligonucleotide crargeted to human microsomal prostaglandin E2 synthase (mPCES-1). The CC human mPCES-1 gene is located on chromosome 9, more specifically to (2) 934.3. The present invention also describes: (1) antisense compounds, (2) antisense compounds, (2) antisense compounds, (3) a method of inhibiting the expression of (4) a method of inhibiting the expression of (5) antisense compounds and (6) antisense compounds (7) a method of treating an animal (7) antisense compounds have cytostatic, (8) antisense com
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ophthalmic,
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGSS-1). The human mPGSS-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds
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                                                                                                                                     Claim 4;
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nilarity 95.0%;
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Pred. No. 1.3e+03;
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; lamunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
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Synthetic.
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16. .20
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25-SEP-2002;

2002US-0413549P

(PHAA ) PHARMACIA CORP

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RESULT 871
ADM15245/c
ID ADM152
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a chimeric antisense oligonucleotide contrared to human microsomal prostaglandin E2 synthase (mpGES-1). The CC human mPGES-1 gene is located on chromosome 9, more specifically to CC 934.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MPGES-1 and CC antisinglammator, candilator, cardiant, neuroprotective, antisinflammatory, neuroprotective, nootropic, antisinflammatory, neuroprotective, nootropic, antisinflammatory, neuroprotective, nootropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's composition injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                         chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpcES-1; mpcES-1 inhibitor; microsomal prostaglandin E2 synthase; mpcES-1; mpcES-1; microsomal prostaglandin E2 synthase; microsomal prostaglandin E2 synthase; antiinflammator; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; disbetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gierse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                          Human mPGES-1
                                                                                                                                                                                                                                                                                                                                                                                                                             01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADM15245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADM15245 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ischemia.
           modified_base
                                                                                                Key
modified_base
                                                                                                                                                                                                           cardiovascular
                                                                                                                                                          Synthetic
                                                                                                                                                                           Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2004-305094/28
                                                                                                                                                                           sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CAAAGTGCTGGGATTACAGG 407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CAAAGTGCTGGGATGACAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                      chimeric antisense oligonucleotide SEQ
                                                                                                                                                                                                            disorder; neurological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NO 266; 132pp; English
                  residues are
                                                                                                                     Location/Qualifiers
/*tag=
                               /notē= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                       *tag=
                                                                   mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     в c;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
                                                                                       σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 18.4; DB 1
Pred. No. 1.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                              disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20;
                                                                                                                                                                                                                                                                                                                                                                                             ID NO:1432
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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Matches
                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mEGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression, (2) a method of inhibiting the expression of mPGES-1 in cells or tissues, and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric having a disease or condition associated with mPGES-1.
                                                                                                                                                                                                                                                                           antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiarthritic, vasotropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; SEQ ID NO 1432; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-305094/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gierse JK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-SEP-2003; 2003WO-US030374
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                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2004028458-A2
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                                                    772
20
                                                                                                               19;
                                                                                                                                       Similarity
                                TTGTATTTTTAGTAGAGATG 791
                                                                                                                                                                                                                               BP; 9
                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2002US-0413549P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "
16. .20
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/mod_base= OTHER
/mote= "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base=
/note= "2'-
                                                                                                                                                                                                                               Ą
                                                                                                                                       1.9%;
                                                                                                                                                                                                                               5 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -O-methocyethyls"
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                                                                                                               <u>.</u>
                                                                                                                                             Pred.
                                                                                                                                                                       Score 18.4;
                                                                                                                  Mismatches
                                                                                                                                                No.
                                                                                                                                             .3e+03
                                                                                                                                                                          DB 1;
                                                                                                                                                                    Length 20;
                                                                                                                     Indels
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e.g.,

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RESULT 872
ADM15422/c
ID ADM15422
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            chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory;
                                                                                       Human mPGES-1 chimeric antisense oligonucleotide SEQ ID
                                                                                                                                                    ADM15422;
                                                                                                                                                                                   standard;
                                                                                                                     (first entry)
                                                                                                                                                                                   DNA;
                                                                                                                                                                                   20
                                                                                           NO:1609
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neuroprotective;

nootropic; antiarthritic;

ophthalmological;

S

Gaps

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                                                                                                                                                                                                                                                   having a sequence comprising 8-30 bp targeted to a nucleic acid mpGES-1 which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiarinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or antisense, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or antisense compound condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition associated with mpGES-1 e.g., inflammation, and iscales or condition associated with mpGES-1 e.g., inflammation, alzheimer's condition associated with mpGES-1 e.g., inflammation and inflammatical condition associated with mpGES-1 e.g., inflammatical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified_base
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                                                                                                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-SEP-2002; 2002US-0413549P
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                                                                                                                                                                                                                                    ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-SEP-2003; 2003WO-US030374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunomodulatory; cardiovascular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2004-305094/28
                                            723
                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ
                                            CTCCTGAGTAGCTGGGACTA 742
                                                                                                                                                                                     BP; 6
                                                                                                                                                                                                                                    immunological,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ij
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NO 1609; 132pp;
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/mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mod_base=
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                                                                                                                                                                                     A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "2'-O-methocyethyls"
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                                                                                                                                                                                                                                 cardiovascular or neurological disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER
                                                                                          0
                                                                                                                Score 18.4;
Pred. No. 1
                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene therapy; inflammation;
                                                                                                              .3e+03;
                                                                                                                                    DB 1; Length 20;
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                                                                                          Gaps
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RESULÉ 873
ADM1468/c
ID ADM146
XX ADM146
XX ADM146
XX ADM146
XX ADM146
XX ADM146
XX Chimer
DE Human
XX Chimer
XX Modifi
FT mod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JUL-2004
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                                                                                                                                                                                                                                                     The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds,
                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-305094/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cardiovascular disorder; neurological disorder; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2004028458-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inflammation, Alzheimer's disease, arthritis, diabetes, cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                      SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                   ID NO 873; 132pp;
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/note= "2'-O-methocyethyls"
16. .20
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residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                      English.
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having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric

antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiarinflammatory, neuroprotective, nootropic, antiarthritic, vasotropi ophthalmological, immunomodulatory and cardiovascular activities, and

vasotropic,

can

be used as mPGES-1 inhibitors and in gene therapy. The antisense componers to be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, ophthalmic, immunological, cardiovascular or neurological disorder.

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RESULT 874
ADM15137/c
IDM 1517/c

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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or ischemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chimeric; antisense oligonucleotide; phosphorothicate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human mPGES-1 chimeric antisense oligonucleotide SEQ ID
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                                                                                                                                                                                                                                                                                                                                            25-SEP-2003;
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                                                                                                                                     WPI; 2004-305094/28
                                                                                                                                                                                                                                                                                        25-SEP-2002; 2002US-0413549P
                                                                                                                                                                                                                                          (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCCAGCTAATTTTTGTATTT 1
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16.20
/*tag= C
/mod_base= OTHER
/mod_base= 07HER
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/mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              residues
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "2'-0-methocyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "phosphorothioate linkages and es are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   σ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 18.4; DB 1; Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               all cytidine
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                                                           e.g.
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XXXDDDDDDDDDDDDDDDDDXXX
                                                 Claim
                                                  4; SEQ
                                                  ID NO 1324; 132pp; English.
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compound

CC targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to CC '934.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC mpGES-1 in cells or tissues; and (3) a method of treating the expression of CC antisense oligonucleotides and antisense compounds have cytostatic, cantidiabetic, immunomodulator, cardiant, neuroprotective, antiarthritic, vasotropic, CC antissiamentory, neuroprotective, nootropic, antiarthritics, and can be used for preparing a composition for treating a disease or CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder. Sequence 20 BP; 4 A; 8 C; 3 G; 5 T; 0 U; 0 Other; The present sequence represents a chimeric antisense oligonucleotide

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Matches
                                                    Query Match
Best Local 9
                     728 GAGTAGCTGGGACTACAGGC 747
                                            19;
20
                                                         Similarity
GAGTAGCTGGGATTACAGGC 1
                                             Conservative
                                                        95.0%;
                                            0; Mismatches
                                                         Score 18.4; DB 1;
Pred. No. 1.3e+03;
                                                                     DB 1;
                                                                    Length 20;
                                               Indels
                                              0,
                                               Gaps
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8

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RESULT 875
ADM15251/c
                                                                              ADM15251 standard; DNA; 20
Human mPGES-1
                           01-JUL-2004
                                                      ADM15251;
                           (first entry)
 chimeric antisense oligonucleotide SEQ
                                                                                ВÞ
    Ħ
    NO:1438
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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.

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Homo
sapiens.
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modified_base
                                                                                                                                                                      Key
modified_base
                                                                                                                                                                                                            Synthetic.
WO2004028458-A2
                                                                                                         modified_base
                                                                                                                                                                         Location/Qualifiers
1. 20
                                                                      /mod_base=
/note= "2'-
                                                           16. .20
                                                                                                                      residues
                                                                                                                        /mod_base= OTHER
/note= "phosphorothioate linkages and
residues are 5-methylcytidines"
                                                                                                *tag=
                         note=
                                                ი
                                                                       -O-methocyethyls
                      -O-methoxyethyls'
                                   OTHER
                                                                                     OTHER
                                                                                                                                        all cytidine
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RESULT 876
ADM13907/c
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9344.3. The present invention also describes: (1) antisense compounds, chaving a sequence comprising 8-30 by targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal mpGES-1 in cells or tissues; and (3) a method of treating an animal chaving a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonuclectides and antisense compounds have cytostatic, antisense oligonuclectides and antisense compounds have cytostatic, antisticiammatory, neuroprotective, nootropic, antiarthritic, vasotropic, antiinfilammatory, neuroprotective, nootropic, antiarthrities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischemia or reperfusion injury, or ophthalmore and condition associated with mpGES-1 e.g., instancial disease.
  Key
                                                                                                                                                                              chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpcES-1; mpcES-1 inhibitor; microsomal prostaglandin E2 synthase; hibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                  ADM13907 standard; DNA; 20
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                                                                                                                  immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                         Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:94.
                                                                                                                                                                                                                                                                                                                                                 01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense compound, having a sequence targeted to a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-SEP-2002; 2002US-0413549P
                                                                                                   cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   encoding
                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             encoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            730 GTAGCTGGGACTACAGGCGC 749
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 1438; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GTAGCTGGGATTACAGGCGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence represents a chimeric antisense oligonucleotide human microsomal prostaglandin E2 synthase (mPGES-1). Th
                                                                                                 injury; opht
lar disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 4 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunological, cardiovascular or neurological disorder.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.9%;
95.0%;
                                                                                                 neurological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                  ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 18.4; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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RESULT 877
ADM13925/c
ID ADM139
XX
AC ADM139
XC
DT 01-JUL
XX

ADM13925; ADM13925

standard;

DNA;

20

ВP

0

01-JUL-2004

(first entry)

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                                                                                                                                     Query Match
Best Local Similarity
                                                                  Matches
                                                                                                                                                                                                                                                                                                                    The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of thibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric
                                                                                                                                                              antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGBS-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alabeimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                     Sequence 20 BP; 4 A; 5 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The CC human mPGES-1 gene is located on chromosome 9, more specifically to CC 934.3. The present invention also describes: (1) antisense compounds, CC mPGES-1, which specifically hybridise with the nucleic acid encoding CC mPGES-1, which specifically hybridise with the nucleic acid encoding CC mPGES-1 in cells or tissues; and (3) a method of inhibiting an animal CC antisense or condition associated with mPGES-1. MPGES-1 and CC antisense or conditions and antisense compounds have cytostatic, CC antisense oligonucleotides and antisense compounds have cytostatic, CC antisinflammatory, neuroprotective, noctropic, antiarthritic, vasotropic, CC ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or CC ophthalmoic, immunological, cardiovascular or neurological disorder.
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l prostaglandin E2 synthase; mPGES-1; mPGES-1
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mBGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encodimpGES-1, which specifically hybridise with the nucleic acid mPGES-1 are
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residues are 5-methylcytidines"
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ADO45368
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ANDO4
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Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (NYCE/) NYCE J W.
(SAND/) SANDRASAGRA F
(TANG/) TANG L.
(AGUI/) AGUILAR D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; 88; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B, PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; adirway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
                                Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g
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                                                                                                                                                                                                                                           Shahabuddin
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23-APR-2002; 2002WO-US013143.
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in S, Lu H, Cong
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ong H;
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Claim 2; SEQ ID NO 734; 174pp; English

The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region codon, coding region, 5' or 3' intron-exon junction, intron or region codon, coding region, 5' or 3'-end or 3'-end of a nucleic acid target cochesen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-6 receptor, CCR1, CCR1, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the CC prevention and/or treatment of a respiratory or lung disease. The CC prevention and/or treatment of a respiratory or lung disease. The CC cran, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The CC respiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A CC receptor(s), and/or asthma and/or levels of adenosine and/or levels of adenosine (c), chronic obstructive pulmonary disease (COPD), chronic obstructive pulmonary disease (COPD), chronic obstructive pulmonary disease (COPD), prevention or This accuse represents an oligonary pulmonary disease in the prevention or the provided respiratory or lung inflammation, bronchitis, airway disease (COPD), and the prevention or the prevention of the prevention or the prevention of the prevention or the prevention or the prevention or the prevention or the prevention of the prevention or the prevention of the prevention or the prevention of the prevention of the preventio bronchoconstriction. This sequence represents an oligonucleotide of the

Sequence 20 BP; 3 A; 6 C; 8 G; 3 T; 0 U; 0 Other;

밁 S Matches Query Match Best Local ( 641 CACCCAGGCTGGAGTGCAGT 660 μ h 1.9%; Similarity 95.0%; CGCCCAGGCTGGAGTGCAGT 20 0 Score 18.4; DB 1; Pred. No. 1.3e+03; Mismatches DB 1; Length Indels 20; 0 Gaps

0

RESULT 880 ADO46436 ADO46436 standard; (first entry) DNA;

Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Ectaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction. 23-APR-2002; 23-APR-2002; 11-MAR-2004. US2004049022-A1 Human oligonucleotide #1802. 25-JUL-2003; 2003US-00627930 2002WO-US013135. 2002WO-US013143.

(NYCE/) (SAND/) (TANG/)

) NYCE J W. ) SANDRASAGRA A. ) TANG L.

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chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-6 chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-7 receptor, interleukin-8 receptor, interleukin-9 receptor, interleukin-9 receptor, interleukin-1 r
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-293804/27.
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(MILL/)
(SHAH/)
(LUHH/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, alrway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region
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                                                                                                                                  Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease;
                                                                                                                                                                                                                                  Human oligonucleotide #629
                                                                                                                                                                                                                                                                               15-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                            ADO45263 standard; DNA;
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                                         lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; chronic obstructive pulmonary disease; COPD; allergic rhinitis;
acute respiratory distress syndrome; pulmonary hypertension;
lung inflammation; bronchitis; airway obstruction; bronchoconstriction
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MILLER S.
SHAHABUDDIN S
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95.0%;
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ong H;
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The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-4 receptor, CCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The CC prevention and/or treatment of a respiratory or lung disease. The CC pene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The CC useful for preventing or treating a respiratory or lung disease. The CC and/or increased levels of, adenosine and/or levels of adenosine of allergic rhinitis, acute respiratory disease. The respiratory or lung disease (COPD), chronic obstructive pulmonary disease (COPD), chronic obstructive pulmonary disease (COPD), allergic rhinitis, acute respiratory bistress syndrome, pulmonary disease and post pulmonary disease are allergic rhinitis, acute respiratory bistress syndrome, pulmonary disease of the pronchactory disease represents an oligonucleotide of the
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                                                             Matches
                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2; SEQ ID NO 629; 174pp; English.
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                                                                                                                          Sequence 20
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                                                                                                                                                                            pronchoconstriction.
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                             542 CTCAGCCTCCCAAGTAGCTG 561
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LU H.
CONG H.
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                                                             l Similarity
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in S, Lu H, Cong
                                                                                                                            вР;
                                                             Conservative
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ong H;
                                                                                                                            5 G; 4 T;
                                                                                                                                                                            sequence represents
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                                                                                             Score 18.4; DB 1; Length 20;
                                                                                Pred.
20
                                                                Mismatches
                                                                              No.
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                                                                                                                                0 Other;
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RESULT 882 ADO45370 ID ADO453 XX

ADO45370 standard; DNA;

20

В₽

ADO45370;

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The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (IL) 4 receptor, interleukin (IL) 5' receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the CC prevention and/or treatment of a respiratory or lung disease. The CC oligonucleotides are useful for reducing or inhibiting expression of a CC gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 C, or PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The CC useful for preventing or treating a respiratory or lung disease. The CC respiratory or lung disease is associated with hyper-responsiveness to a receptor(s), and/or asthma and/or lung allergies associated with a receptor(s), and/or asthma and/or lung allergies associated with constituents of a stream and/or lung allergy, asthma, impeded respiration, CC cystic fibrosis (CP), chronic obstructive pulmonary disease (CDD), chronic constructive pulmonary disease (CDD).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; SEQ ID NO 736; 174pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
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23-APR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human oligonucleotide #736
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ng H;
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ronchitis, airway obstruction or represents an oligonucleotide of
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RESULT 883
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Best Local (
The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL)-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The oligonucleotides are useful for reducing or inhibiting expression of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
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                                                                                                                                                                  Claim 2;
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(TANG/)
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                                                                                                                                                               SEQ ID NO 623; 174pp; English
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Pred. No. 1.3e+03
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                                                                                                                                                                                                             respiratory disease e.g.
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RESULT 884
ADO45258
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                               Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; CCRM; tryptase a; tryptase b; PD24 A; PD24 B; PD24 C; PD24 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
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(SAND/)
(TANG/)
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                                                                                                                                                                                                                                                                            23-APR-2002;
Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g.
                                           WPI; 2004-293804/27
                                                                        Nyce JW, Sandrasagra
Shahabuddin S, Lu H,
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TANG L.
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SHAHABUDDIN S
                                                                                                                                                                                AGUILAR D.
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Cong
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)ng H;
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Pred. No. 1.3e+03;
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codon, coding region, 5' or 3' intron-exon junction, intron or region cotion, coding region, 5' or 3' intron-exon junction, intron or region come with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (II) 4 receptor, interleukin (II) 5- receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CCR tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s). For the coligonuclectides are useful for reducing or lung disease. The coligonuclectides are useful for reducing or inhibiting expression of a gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CC CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonuclectides are useful for preventing or treating a respiratory or lung disease. The CC espiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A CC receptor(s), and/or asthma and/or lung allergies associated with cC inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary disease (COPD), cronic obstructive pulmonary disease (COPD), coligonucleotide of the bronchoconstriction. This sequence represents an oligonucleotide of the
Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; SEQ
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Matches Query Match Best Local S Sequence 20 873 ACAGGCGTGAGCCACCACGC 892 **\_** 19; Similarity ACAGGCGTGGGCCACCACGC 20 B₽; Conservative 4 ð 1.9%; ω C; 7 G; 1 T; <u>.</u>. Score 18.4; Pred. No. 1 Mismatches 0 u; 1.3e+03 BG 0 Other; 1; Length Indels 20; 0 Gaps 0

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PRESENTATION OF THE PRESEN RESULT 885 Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Botaxin-1; RANTES; MCP4; CD23; ICAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; alivary inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; acute respirator; bronchotonstriction. 23-APR-2002; 23-APR-2002; 25-JUL-2003; 2003US-00627930 11-MAR-2004. US2004049022-A1 Homo sapiens. Human oligonucleotide #1817. 15-JUL-2004 ADO46451; ADO46451 standard; DNA; 20 2002WO-US013143 2002WO-US013135 (first entry)

(NYCE/) NYCE J

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RESULT 886
ADO46479
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Best Local S
Matches 19
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Shahabuddin
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                  lung disease; hyper-responsiveness; adenosine; adenosine A receptor;
asthma; lung allergy; inflammation; inflammatory disease;
airway inflammation; allergy; impeded respiration; cystic fibrosis; CF.
                                                                                     Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease;
                                                                                                                                                                               Human oligonucleotide #1845.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction obronchoconstriction. This sequence represents an oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; SEQ ID NO 1818; 174pp; English.
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(MILL/)
(SHAH/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20
chronic obstructive pulmonary
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TANG L.
AGUILAR D.
MILLER S.
SHAHABUDDIN
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in S, Lu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 6 A;
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ong H;
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disease; COPD; allergic
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                                                                   Matches
                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
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                                                                                                                                    Sequence 20 BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
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CONG H.
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                     CTGGTCTCAAACTCCTGACC 1134
CTGGTCTCAAACTCCTGAGC
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                                                                   Conservative
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2002WO-US013143.
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                                                                                                                                                                                     This sequence represents an oligonucleotide
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ng H;
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                                                                                   Score 18.4; DB 1;
Pred. No. 1.3e+03;
20
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RESULT 887 ADO45320

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chosen from a gene encoding interleukin (IL) 4 receptor, interleukin (IL) crossen from a gene encoding interleukin (IL) 4 receptor, CR1, CR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CT tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention calso relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The C cligonucleotides are useful for reducing or inhibiting expression of a gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, C CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, C CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, C C CR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, C C crasses b, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The crespiratory or lung disease to add/or increased levels of, adenosine and/or levels of adenosine A C inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammator, allergy, asthma, impeded respiration, c cystic fibrosis (CP), chronic obstructive pulmonary disease (CPD),
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23-APR-2002; 2002WO-US013143.
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                                 allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction o
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AGUILAR D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sandrasagra
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represents an oligonucleotide of the
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RESULT 888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; ss; interleukin-4 receptor; II-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Ectaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (II)-4 receptor, interleukin (I-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDB4 A, PDB4 B, PDB4 C or PDB4 D. The invention also relates to a method of screening a candidate compound that binds tone or more nucleic acid target(s) or expressed product(s), for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 2
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                                                                                                                                                                                                                                                                                                           Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-293804/27.
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23-APR-2002;
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                                                                                                                                                                                                                                           Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (SHAH/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NYCE J W.
SANDRASAGRA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SHAHABUDDIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TANG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TU H.
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                                                                                                                                                                                                                                           SEQ ID NO 724; 174pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sandrasagra
in S, Lu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  interleukin-4 receptor; IL-4; interleukin-5 receptor;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      a A,
Cong
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ong H;
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Pred. No. 1
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Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             prevention and/or treatment of a respiratory or lung disease. The oligonucleotides are useful for reducing or inhibiting expression of a gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The respiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A receptor(s), and/or asthma and/or lung allergies associated with inflammation or an inflammation, disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, cystic fibrosis (CP), chronic obstructive pulmonary disease (COPD),
                                                                                                                                                                                                                                                                                                                                                                                                                           Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; CCMx tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; CPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
               WPI; 2004-293804/27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human oligonucleotide #1811.
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                                            Shahabuddin S,
                                                               Nусе JW,
                                                                                                                                                                                                                                                 23-APR-2002; 2002WO-US013135.
23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                  US2004049022-A1.
                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                             TANG L.
AGUILAR D.
MILLER S.
                                                                                                CONG H.
                                                                                                                                                                                               SANDRASAGRA A.
                                                                                                                                                                                                                  NYCE J W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                             SHAHABUDDIN S
                                           Sandrasagra A, Ta
in S, Lu H, Cong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 5 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.9%;
                                            Tang L, ong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВP
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Pred. No. 1.3e+03;
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                                                            Aguilar D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 20;
                                                               Miller
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The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region complete with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-5 receptor, CCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PD84 A, PD84 B, DD84 C or PD84 D. The invention collection and/or treatment of a respiratory or lung disease. The collection and/or treatment of a respiratory or lung disease. The collisonucleotides are useful for reducing or inhibiting expression of a collisonucleotides are useful for receptor, interleukin-5 receptor, cCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, ctryptase b, PD84 A, PD84 B, PD84 C, or PD84 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The cceptor(s), and/or asthma and/or lung allergies associated with receptor(s), and/or asthma and/or lung allergies associated with crecaptor(s), and/or asthma and/or lung allergies associated with collection or an inflammatory disease. The respiratory or lung disease collection stomation, lung disease collection, constructive pulmonary disease (COPD), chronic obstructive pulmonary disease (COPD), collection of the properties of action or this accuracy or some pulmonary disease collection or inflammation, bronchitis, across syndrome, pulmonary collection or the properties of the collection or the pulmonary disease collection or the properties of the collection or the pulmonary disease collection or the properties of the collection or the pulmonary disease colle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; SEQ ID NO 1812; 174pp; English
                                                                                                             pronchoconstriction. This sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            e.g.
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밁 ঠ Matches Query Match Best Local Similarity ADO45319 standard; DNA; 20 199 19; 1 ATGTTGGCCAGGCTGGTCTC 20 ATGTTGGTCAGGCTGGTCTC 218 Conservative 1.9%; <u>.</u>. Score 18.4; DB 1; Pred. No. 1.3e+03; Mismatches DB 1; Length ۲. Indels 20 0; Gaps

0

Sequence 20 BP; 2 A; 5 C; 7 G; 6 T; 0 U; 0 Other;

Human; ss; interleukin-4 receptor; II-4; interleukin-5 receptor; II-5; CCR1; CCR3; Eotaxin-1; RANTES; MCCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PD54 A; PD54 B; PD54 C; PD54 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; allergy; inflammatory disease; corpo; allergic rhinitis; cronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; 23-APR-2002; 2002WO-US013135. 23-APR-2002; 2002WO-US013143. US2004049022-A1 Human oligonucleotide #685 15-JUL-2004 (first entry) 25-JUL-2003; 2003US-00627930. Homo sapiens inflammation; bronchitis; airway obstruction; bronchoconstriction

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-6 consent from a gene encoding interleukin (II)-7 receptor, interleukin (II)-7 consent from a gene encoding interleukin-8 pDB4 Cor pDB4 D. The invention consent from and/or treatment of a respiratory or lung disease. The consent from and/or treatment of a respiratory or lung disease. The consent from and/or treatment of a respiratory or lung disease in consent from a respiratory or lung disease. The consent from preventing or treating a respiratory or lung disease. The consent from a find for preventing a respiratory or lung disease. The consent from a find for and/or lung allergies associated with hyper-responsiveness to inflammation or an inflammatory disease. The respiratory or lung disease is associated with hyper-responsiveness for inflammation or an inflammatory disease. The respiratory or lung disease is consent from a firm inflammation, allergy, asthma, impeded respiration, constitution in the consent from a constitutive pulmonary disease (CPPD), consider from a constitution of the constitutive pulmonary disease (CPPD).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
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(AGUI/)
(MILL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target
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Human; 88; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease;
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       bronchoconstriction.
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                                                                                               Human oligonucleotide #630.
                                                                                                                                                                                            ADO45264 standard; DNA; 20 BP
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MILLER S.
SHAHABUDDIN
LU H.
CONG H.
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19; Conserv
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SANDRASAGRA
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airway inflammation; allergy; impeded respiration; cystic fibrosis; CF chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction
Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
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(SAND/)
(TANG/)
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MILLER S.
SHAHABUDDIN
LU H.
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TANG L.
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in S, Lu H, Cong
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2002WO-US013143.
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Claim 2; SEQ ID NO 630; 174pp; English.

WWW. Social control of the control o

asthma.

The invention relates to oligonucleotides anti-sense to an initiation CC codon, coding region, 5' or 3' intron-exon junction, intron or region CC with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target CC chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL)-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The CC oligonucleotides are useful for reducing or inhibiting expression of a CC gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CC CTR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC tryptase b, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC tryptase b, DE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC tryptase b, DE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC tryptase b, DE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC tryptase b, DE5 B, DE5

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Sequence 20 BP; 3 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
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                                   Query Match
Best Local
              722
                                     Similarity
              CCTCCTGAGTAGCTGGGACT 741
                             Conservative
                                    1.9%;
95.0%;
                             o
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                                     Score 18.4;
Pred. No. 1
20
                              Mismatches
                                     .3e+03
                                             DB 1;
                                             Length
                                Indels
                                             20;
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RESULT 892
ADO45367
       CC The invention relates to oligonucleotides anti-sense to an initiation CC with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target CC chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-5 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to come or more nucleic acid target(s) or expressed product(s), for the CC prevention and/or treatment of a respiratory or lung disease. The CC oligonucleotides are useful for reducing or inhibiting expression of a CC gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CC CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are cuseful for preventing or treating a respiratory or lung disease. The CC compliance of levels of, adenosine and/or levels of adenosine A CC craft and/or asthma and/or lung allergies associated with CC inflammation or an inflammatory disease. The respiratory or lung disease conforms from airway inflammation, allergy, asthma, impeded respiration, CC cystic fibrosis (CF), chronic obstructive pulmonary disease (CDPD), and control of the co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel single or multiple target oligonucleotide anti-sense to e.g. coninitiation codon, intron of respiratory disease-relevant gene e.g. CC RANTES, MCP4, useful for prophylaxis or treating respiratory disease
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23-APR-2002; 2002WO-US013143.
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AGUILAR D.
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SHAHABUDDIN S.
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in S, Lu H, Cong
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acute
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respiratory distress
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syndrome,
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RESULT 893
AD013029
ID AD0130
XX AD0130
XX Single
XX Geneti
XX Geneti
XX W02004
XX C2-APF
XX W70-OCT
XX W70-OCT
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                                                                                                The invention relates to a method of designing primers for simultaneous camplification of target DNA fragments in a single multiplex polymerase contain reaction by aligning a first primer and a second primer. The method comprises: (a) aligning a first primer and a second primer; and (b) comprises: (a) aligning a first primer and a second primer; and (b) contain four or more bases that are perfectly matching to the 3' end cos not contain seven or more bases that are perfectly matching to the 3' end cos not contain seven or more bases that are perfectly matching coscept one mismatch to the 3' end sequence of the first primer at its 3' end does not contain six or contain six or more bases that are perfectly matching coscept one mismatch to the 3' end sequence anywhere of the first primer or the second primer, and the first primer at its 3' end does not contain six or contain eleven or more bases that are perfectly matching except cone mismatch to a sequence anywhere of the first primer at its 3' end contain six or more bases that are perfectly matching except cone mismatch to a sequence anywhere of the first primer or the second contain eleven or more bases that are perfectly matching except cone mismatch to a sequence anywhere of the first primer or the second contain eleven or more bases that are perfectly matching except cone mismatch to a sequence anywhere of the first primer or the second contain eleven or more bases that are perfectly matching except cone mismatch to a sequence anywhere of the first primer or the second contain eleven or more bases that are perfectly matching except cone mismatch to a sequence anywhere of the first primer or the second contain eleven or more bases that are perfectly matching enemals. The second contain eleven or more bases that are perfectly matching enemals and the first primer or the second contain eleven or more bases that are perfectly matching enemals.
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single multiplex polymerase chain reaction; multifactorial disease;
genetic alteration; pharmacogenetic reaction; genotyping; polymorph
alterations, the studies in pharmacogenetic reactions, the genotyping genetic polymorphisms in a large population, the gene expression profiling in various samples and high throughput genotyping technolog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase chain reaction, for high throughput
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 44; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             multiplex DNA sequence amplification, comprises aligning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-340914/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2004033649-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Single multiplex PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYNE-) UNIV NEW JERSEY MEDICINE & DENTISTRY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             expression
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 Similarity
19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         μ.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 2 A; 6 C; 7 G; 5 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2002US-0417009P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ion; pharmacogenetic reaction; genotyping; polymorphism;
profiling.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВÞ
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1;
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technologies

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RESULT 894
ADN58838/c
ID ADN588
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RESULT 895
ADP70377
ID ADP703
XX
AC ADP703
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Matches
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Best Local (
                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This
                                                                                                                                                                                                            The invention relates to a compound targeted to a nucleic acid molecule encoding B7H, where the compound specifically hybridises with the nucleic acid molecule encoding B7H and inhibits the expression of B7H. The compound is useful for modulating the expression of B7H. It is also useful for diagnosing or treating diseases associated with expression of B7H, e.g., an autoimmune disease. The present sequence represents a human
                                                                                                                                                                                                                                                                                                                 New compound targeted to a nucleic acid molecule encoding B7H and inhibits expression of B7H, useful for modulating the expression or for diagnosing or treating, e.g. autoimmune disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2004102398-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human B7H antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-NOV-2002; 2002US-00303420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADN58838 standard;
                                                                                                                                                                             Sequence 20 BP; 4 A; 5 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                              Example 15; SEQ ID NO 89; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                 23-NOV-2002; 2002US-00303420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                  ADP70377 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                 2004-399728/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence corresponds to an example of a primer of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   484 AGTGGTGTGATCACAGCTCA 503
                                                                                                            870
                                                                                                                                  al Similarity
19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19;
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                                                                                                           ATTACAGGCGTGAGCCACCA 889
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                                                                                                                                  1.9%;
ilarity 95.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
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                                   BP.
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Pred. No. 1
                                                                                                                                              Score 18.4;
Pred. No. 1.
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                                                                                                                                   Mismatches
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                                                                                                                                               1.3e+03
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                                                                                                                                                         DB 1;
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                                                                                                                                                       Length 20;
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Omo

sapiens

Human

Ephrin-B2 DNA antisense oligonucleotide

26-AUG-2004

(first entry)

Human; Ephrin-B2; ss; antisense oligonucleotide; phosphorothioate linkage; 2'-O-methoxyethyl suga 5-methylcytosine; hyperproliferative disorder; c

sugar moiety; cancer;

cytostatic.

10-DEC-2002; 2002US-00316516

10-JUN-2004. US2004110150-A1

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RESULT 896
ADP26815
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel peptide comprising a TRG1-20 sequence capable of recognising and inducing the human leukocyte antigen (HLA)-B52 or HLA-B62 constraint property of a cytotoxic T-cell (CTL) or a peptide comprising a TRG2-41 sequence capable of recognising and inducing the HLA-B52 of a CTL. The peptide of the invention demonstrates cytostatic activity and may be useful for inducing a cytotoxic T-cell in order to treat cancer, preferably epithelial cancer, more preferably lung cancer, stomach cancer, colon cancer, prostatic cancer and/or melanoma. The treatment may comprise the use of a vaccine. The current sequence is that of the PCR primer 4 of the invention which was used to analyse human testin-related gene (TRG) expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel tumor antigens TRG1-20 and TRG2-41 capable of recognizing inducing human leukocyte antigen B52 or B62 constraint property cytotoxic T lymphocyte, useful for treating cancer e.g., colon of the co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human leukocyte antigen; HLA-B52;
TRG2-41; TRG1-20; cytostatic; epit
prostate; melanoma; vaccine; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-AUG-2004 (first entry)
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ADP26815 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                   536 TCCTGCCTCAGCCTCCCAAG
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             352; HLA-B62; cytotoxic T-cell; CTL; epithelial cancer; lung; stomach; colon; numan; testin-related gene; ss; PCR; primer.
                                                                                                                                                                                                                                                      20
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., colon cancer,
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Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a compound targeted to a nucleic acid molecule encoding the human Ephrin-B2 polypeptide. The compound is an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibits expression of the polypeptide. The antisense oligonucleotide comprises at least one modified internucleoside linkage i.e. a phosphorothicate linkage, at least one modified sugar moiety, preferably a 2'-O-methoxyethyl sugar moiety, or at least one modified nucleobase comprising a 5-methylcytosine. The antisense compounds are useful for modulating the expression of the human Ephrin-B2 polypeptide and in preparation of a composition for treating hyperproliferative disorders, e.g. cancer. This sequence represents an antisense oligonucleotide targeted to DNA encoding the human Ephrin-B2 polypeptide of the
                                                                                                                                                                                                                                                                                                    Human; Ephrin-B2; ss; antisense oligonucleotide; phosphorothicate linkage; 2'-O-methoxyethyl sugar moiety; 5-methylcytosine; hyperproliferative disorder; cancer; cy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 15; SEQ ID NO 64; 69pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New oligonucleotide compound that inhibits expression of Ephrin-B2.
           New oligonucleotide compound that inhibits expression of Ephrin-B2, useful for preparing a composition for treating hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                Human Ephrin-B2 DNA antisense oligonucleotide target region
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADP26872 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-DEC-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
                                                                                                                                                     10-DEC-2002; 2002US-00316516
                                                                                                                                                                                                                                            US2004110150-A1
                                                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                                                                   10-DEC-2002; 2002US-00316516
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                                                          2004-440339/41.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GCTAGGATTACAGGCGTGAG 20
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preparing a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 C; 8 G; 4 T; 0 U; 0 Other;
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Pred. No. 1.3e+03;

    Length 20;

                                                                                                                                                                                                                                                                                                        cancer; cytostatic
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encoding the human Ephrin-B2 polypeptide. The compound is an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibits expression of the polypeptide. The antisense oligonucleotide comprises at least one modified internucleoside linkage i.e. a phosphorothioate linkage, at least one modified sugar moiety, preferable a 2'-O-methoxyethyl sugar moiety, or at least one modified nucleobase comporising a 5-methylcytosine. The antisense compounds are useful for modulating the expression of the human Ephrin-B2 polypeptide and in preparation of a composition for treating hyperproliferative disorders, e.g. cancer. This sequence represents a human Ephrin-B2 DNA antisense oligonucleotide target region of the invention.
Sequence 20 BP; 4 A; 8 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                      Example 15; SEQ ID NO 121; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                      to a
                                                                                                                                                                                                                                                                                                                                                   compound targeted to a nucleic acid molecule
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        864 GCTGGGATTACAGGCGTGAG 883
20
                           19; Conserv
                            Conservative
                                  95.0%;
                                         1.9%;
                           0
                                   Pred. No.
                                         Score 18.4;
                            Mismatches
                                   1.3e+03
                                          DB 1;
                                          Length 20;
                            Indels
                            0
                           Gaps
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RESULT 898
AD071539/c
ID AD0715
XX AD0715
XX AD0715
XX Done m
KW Chromo
KW Lumbar
XX Homo E
XX Homo E
XX W21-NOV
XX 21-NOV
XX 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-NOV-2003; 2003WO-GB005055.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-JUN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Forward primer for SNPs in exon
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VINU (-BAYU)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                osteoporosis;
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The specification describes a method for assessing bone mineral density (BMD) in an individual. The method comprises using a chloride channel 7 (Clcn7) gene marker. The Clcn7 gene maps to chromosome 16p13 and comprises 25 exons. This polymorphic marker is a single nucleotide polymorphism (SNP)in position 14476 situated in intron 8, position 19233 situated in exon 15, position 19240 situated in exon 15, position 39699 situated in exon 1, or position 39705 situated in exon 1. The polymorphic marker may also be a tandem repeat marker which is the 50 bp repeat polymorphic marker which is in linkage disequilibrium with it. The method of the Claim Assessing bone mineral density (BMD) in an individual, useful for treating the individual to prevent or reduce the onset of osteoporosis, comprises using a chloride channel 7 (Clcn7) gene marker. 24; Page 25; 51pp; English.

WPI; 2004-420640/39.

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                                                                                                                                               The invention relates to a novel method for identifying a subject at risk of breast cancer which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a risk of breast cancer, as well as therapeutic and prophylactic treatments that specifically target breast cancer, such as gene therapy. The current sequence is that of an Extend primer of the invention which was used to genotype single nucleotide polymorphisms within human glycoprotein VI (platelet) (GP6; GPIV;GPVI) DNA which is located at chromosomal position 19q13.4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      breast cancer; cytostatic; gene therapy; GPG, GPIV; GPVI; chromosome 19q13.4; ss;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying a subject at risk of breast cancer by detecting the presence or absence of one or more nucleotide polymorphic variations, useful for diagnosing, preventing and/or treating breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-NOV-2002; 2002US-0429136P.
24-JUL-2003; 2003US-0490234P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; Page 82; 286pp; English.
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                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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1.9%;
                                                                                         7 C; 5
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                                                                                             G; 4 T; 0 U;
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Pred. No. 1
   Score 18.4;
Pred. No. 1
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                                                                                                0 Other;
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                                 DB 1;
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Similarity

ADG70429 ID ADG7 XX AC ADG7 XX

ADG70429 standard; DNA;

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ВP

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ADG70429

RESULT 901

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ARTSULT 900
ARTS2349/c
ID AATS23
XX
AAATS23
AC AAATS23
XX
DT 11-JUN
XX Bubble
KW PCR; p
XX US5597
XX UNITO6
XX UNITO
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                                                                                                                                                                                                                                                                     (AAT62346). The method involves ligating a double stranded DNA structure with a non-complementary region, a 'bubble', in the centre (e.g. see AAT62343-4), to restriction digested fragments of regions containing IRES. The ligation results in a double stranded DNA molecule containing at least one 'bubble' at either end. After denaturing the structure, amplification of the IRE-containing region proceeds by PCR using primers targeted to the IRE sequence (e.g. AAT62347-50) and to the sequence in the 'bubble' region (e.g. see AAT62345). The primer presented here binds to nucleotides 216-236 of the Alu-J polymorphic repeat sequence. The method can be used to detect the presence or absence of a chromosomal aberration e.g. in a genetic disorder, in a test organism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bubble; interspersed repetitive element; ligation; annealing; PCR; polymerase chain reaction; amplification; chromosomal abs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amplification of nucleic acid having interspersed repetitive element - using bubble oligo:nucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            genetic disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Primer Alu-J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT62349;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to the amplification of region of DNA containing interspersed repetitive elements (IRB) such as the Alu repeat sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Col 17; 16pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Munroe
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                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US5597694-A
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21
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                                                                                                            l Similarity
19; Conserv
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                                                      ATCACAGCTCACTGCAGCCT 512
                                                                                                                                                                                                                              21
   ATCACGGCTCACTGCAGCCT 2
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                                                                                                               Conservative
                                                                                                                                                                                                                              BP; 4 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                              4 C; 9 G; 4 T;
                                                                                                                                       1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21
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                                                                                                               0
                                                                                                                                          Score 18.4;
Pred. No. 1.
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                                                                                                                     Mismatches
                                                                                                                                                                                                                                 ou;
                                                                                                                                       .4e+03
                                                                                                                                                                                                                                    0 Other;
                                                                                                                                                                        DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                        Gaps
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11-MAR-2004

(first entry)

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PART X BARAN SALAN BARAN BARAN
                                                                                                                                                                                                                         RESULT 902
ADG70430/c
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   BXBXBX8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel isolated or recombinant nucleic acid compensing an ANGE, CLLD8 or CLLD7 mRNA, or ANGE-CLLD8, ANGE-CLLD8, CLLD7, antiasthmatic, dermatological, antipyretic, and antiinflammatory. The nucleic acids of the invention may be used in gene therapy to treat disorders. The nucleic acid sequences are useful for screening agents that inhibit or enhance activity of an ANGE, CLLD8, or CLLD7 gene. The agent or antibody is useful for treating agents that inhibit or enhance activity of an ANGE, CLLD8, or CLLD7 gene. The agent or antibody is useful for treating assay detecting or measuring the polypoptide in the sample. The host cell is useful for producing, regulating and analyzing the polypoptide. The splice variant of ANGE, CLLD8, or CLLD7 is useful for producing the severity, or predisposition to a disease. This polynucleotide sequence represents an REN-34 SNP binding colling to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New nucleic acid sequence comprising an ANGE, CLLD8 or CLLD7 mRNA, or their hybrid, useful for screening agents for treating IgE-mediated diseases, e.g. asthma, atopy, hay fever, eczema, atopic dermatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-201405/19.
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21-JUN-2001; 2001GB-00015212.
21-JUN-2001; 2001GB-00015213.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANGE; CLLD8; CLLD7; ANGE-CLLD8; ANGE-CLLD7, CLLD7-CLLD8; ANGE-CLLD8-CLLD7; antiallergic; antiasthmatic; dermatological; antipyretic; antiinflammatory; gene therapy; IgE-mediated disease; REN-34; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REN-34 SNP binding area oligo #3
REN-34 SNP binding area oligo #4.
                                                               11-MAR-2004
                                                                                                                                                                                              ADG70430 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 429; 429pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               allergic rhinitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zhang
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                                                                                                                                                                                                                                                                                                                                                                                                                        589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               relating to the invention.
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                                                                                                                                                                                                                                                                                                                                                        CTCTGCCTCCTGGGTTCAAG 20
                                                                                                                                                                                                                                                                                                                                                                                                                        CTCTGCCTCCCGGGTTCAAG 704
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 2 A; 8 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cookson
                                                                                                                                                                                              21
                                                                                                                                                                                              BP
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 21;
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15-JUL-2004

(first entry)

Single multiplex PCR primer #2375.

ss; primer; simultaneous amplification;
single multiplex polymerase chain reaction; multifactorial disease;

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RESULT 903
                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC Sequence comprising an ANGE, CLLD8 or CLLD7 mRNA, or ANGE-CLLD8, ANGE-CLLD8, CLLD7, CLLD8, or ANGE-CLLD8 CLLD8 or CLLD7 mRNA sequence, its CC complement, homologue or fragment. The novel nucleic acid sequences have the following activities: antiallergic, antiasthmatic, dermatological, CC antipyretic, and antiinflammatory. The nucleic acids of the invention may be used in gene therapy to treat disorders. The nucleic acid sequences care useful for screening agents that inhibit or enhance activity of an ANGE, CLLD8 or CLLD7 gene. The agent or antibody is useful for treating CC IgE-mediated diseases, such as asthma, atopy, hay fever, eczema, atopic cuseful in an assay detecting or measuring the polypeptide in the sample. CC Useful in an assay detecting or measuring the polypeptide in the sample. CC Olypeptide. The splice variant of ANGE, CLLD8, or CLLD7 is useful for CC diagnosing an IgE-mediated disease, atopy, a form of atopic disease or con-atopic asthma, or predicting the severity, or predisposition to a CC disease. This polynucleotide sequence represents an REN-34 SNP binding conjugation of the invention.
                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-JUN-2001;
21-JUN-2001;
21-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New nucleic acid sequence comprising an ANGE, CLLD8 or CLLD7 mRNA, or their hybrid, useful for screening agents for treating IgE-mediated diseases, e.g. asthma, atopy, hay fever, eczema, atopic dermatitis, or allergic rhinitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ANGE; CLLD8; CLLD7; ANGE-CLLD8; ANGE-CLLD7; CLLD7-CLLD8; ANGE-CLLD8-CLLD7; antiallergic; antiasthmatic; dermatological; antipyretic; antiinflammatory; gene therapy; IgE-mediated disease;
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 6 A; 5 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel isolated or recombinant nucleic acid sequence comprising an ANGE, CLLD8 or CLLD7 mRNA, or ANGE-CLLD8, ANGE-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 429; 429pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-201405/19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS INNOVATIONS LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-JUN-2002; 2002WO-GB002859
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ADO13003
                                                                                                                                                                                                                       589
                                                                                                                                                                                                                                                                                     19;
                                                                                                                                                            21
   standard; DNA;
                                                                                                                                                                                                                    CTCTGCCTCCCGGGTTCAAG 704
                                                                                                                                                            CTCTGCCTCCTGGGTTCAAG 2
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                                                                                                                                                                                                                                                                                     Conservative
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2001GB-00015212.
2001GB-00015213.
                                                                                                                                                                                                                                                                                                               95.0%;
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                                                                                                                                                                                                                                                                                                                                                   1.9%;
   21 BP
                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                      Pred. No.
                                                                                                                                                                                                                                                                                                                                             Score 18.4;
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                                                                                                                                                                                                                                                                                                                      1.4e+03
                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                Length 21;
                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                     0:
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RESULT 904
AAF84350/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a method of designing primers for simultaneous camplification of target DNA fragments in a single multiplex polymerase chain reaction by aligning a first primer and a second primer. The method comprises: (a) aligning a first primer and a second primer; and (b) comprises the first primer where the first primer at its 3' end does not contain four or more bases that are perfectly matching to the 3' end cost and does not contain seven or more bases that are perfectly matching comprise of the first primer at its 3' end does not contain seven or more bases that are perfectly matching comprise or the second primer, the first primer or the second primer, the first primer or the second primer, the first primer or the second primer at its 3' end does not contain six or comprise that are perfectly matching to a sequence anywhere of the first primer at its 3' end does not contain eleven or more bases that are perfectly matching except one mismatch to a sequence anywhere of the first primer at its 3' end does not contain eleven or more bases that are perfectly matching except come mismatch to a sequence anywhere of the first primer or the second complification of target DNA fragments in a single multiplex polymerase contain reaction. It is also useful in the identification of multiple genes crelated to multifactorial diseases, the genome-scale detection of genetic polymorphisms in a large population, the gene expression contains in a large population, the gene expression contains in a single multiple genes or this sequence corresponds to an example of a primer of the invention.
Matches
                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           genetic alteration; pharmacogenetic reaction; genotyping; polymorphism;
gene expression profiling.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase chain reaction, for high throughput multiplex DNA sequence amplification, comprises aligning two primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-OCT-2002; 2002US-0417009P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-340914/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This sequence corresponds to an example of a primer
                                                      Gene polymorphism; drug-metabolising enzyme; PCR primer; CYP2C18i;
                                                                                             Human CYP2C18i PCR primer #6
                                                                                                                                   20-JUN-2001
                                                                                                                                                                                                              AAF84350 standard;
                                                                                                                                                                                                                                                                                                                                                 188
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                                                                                                                                                                                                                                                                                                                                                                                      19;
                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                 GGAGTTTCTCCATGTTGGTC 207
                                                                                                                                                                                                                                                                                                             GGGGTTTCTCCATGTTGGTC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 1 A;
                                                                                                                                   (first entry)
                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                      1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5 C; 7 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                        Score 18.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                          4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                        Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                        0,
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                                                            88.
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                                                                                                                                                                                                     Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                             gene polymorphisms of drug-metabolising enzyme genes. The kit contains a polymerase chain reaction (PCR) buffer solution containing DNA polymerase and NTP, a normal forward primer, a mutated forward primer, a reverse primer and a fluorescence-labelling probe. The method involves carrying out PCR on sample DNA, containing a drug-metabolising enzyme gene, together with PCR buffer, the normal forward primer, the reverse primer and the fluorescence-labelling probe (step A); and carrying out PCR on the sample DNA together with PCR buffer, the mutated forward primer, the reverse primer and the fluorescence-labelling probe (step B), and a step of comparing the result of step a with that of step B). The present sequence is a primer for human CYP2C18i, which was used to illustrate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JP2001017185-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detection of gene polymorphism of drug-metabolizing enzymes useful for diagnosis and testing comprises carrying out polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-MAR-1999;
06-MAY-1999;
                                                                                                                                                                                                                                                                   Sequence 22 BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a kit and method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-285409/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 13; 27pp; Japanese.
20-MAY-1998
                                AAV06198;
                                                               AAV06198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SAKA ) OTSUKA
                                                                                                                                                                       863 TGCTGGGATTACAGGCGTGA 882
                                                                                                                                            20
                                                                                                                                                                                                          19;
                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                   invention
                                                             standard;
                                                                                                                                          TGCTGGGATTACAGGCATGA 1
                                                                                                                                                                                                          Conservative
 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99JP-00076592.
99JP-00125918.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99JP-00351610
                                                               DNA;
                                                                                                                                                                                                                     1.9%;
                                                                                                                                                                                                                                                                    8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CO LID
                                                                                                                                                                                                                                                                    C; 3 G; 5 T; 0 U; 0 Other;
                                                               23
                                                                                                                                                                                                 0;
                                                                                                                                                                                                                        Score 18.4;
Pred. No. 1
                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                        .4e+03
                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                     Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for the detection of
nes. The kit contains a
                                                                                                                                                                                                          Indels
                                                                                                                                                                                                          0,
                                                                                                                                                                                                          Gaps
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RESULT 905
AAV06198/c
Short tandem repeat loci; D3S1539; D4S2368; D5S818; D7S820; D9S930; D10S1239; D13S317; D14S118; D14S562; D16S490; D16S539; D16S1239; D15S1298; D19S1299; D19
                                                                                                                                        15-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           multiplex amplification reaction; MAR; allele;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Primer used when one of the loci in the MAR set is D22S683
                                                  (PROM-) PROMEGA CORF
                                                                                                                                                                                                                                                                                                                                         23-OCT-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9739138-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linkage
                                                                                                                                                                                                                                         15-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              map;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              identification; disease gene; PCR primer; amplify; ss
                                                                                                                                             96US-00632575
                                                                                                                                                                                                                                         97WO-US006293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                detection; genetic marker;
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D16S753;

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AAA47246/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CC Primers AAV06168-228 are used in a novel method for simultaneously CC determining the alleles present in short tandem repeat loci from one or CC more DNA samples. The DNA sample to be analysed has a set of at least CC consisting of D3S1539, D4S2368, D5S818, D7S820, D9S930, D10S1239, CC D13S17, D14S118, D14S548, D14S562, D16S490, D16S539, D16S753, D17S1298, CC D17S1299, D19S253, D2OS481, D2S2633, HUMCSF1DP, HUMFDAX, HUMFIO1, CC HUMFBSFPS, HUMF13A01, HUMBFXIII, HUMLIPOL and HUMWFDAX, Lander repeat loci the DNA sample to be analysed has a set of three short tandem repeat loci CC which can be amplified together, where the set of loci is selected from CC the DNA sample to be sets: (1) D3S139, D19S253, D13S17; (2) D10S1239, CC D9S930, D2OS481; (3) D10S1239, D4S3368, D2OS481; (3) D10S1239, D4S3317, The CC loci are co-amplified in a multiplex amplification reaction (MAR), where the product of the reaction is a mixture of amplified alleles from each CC of the co-amplified loci in the set. The amplified alleles in the mixture CC are evaluated to determine the alleles present at each of the loci analysed in the set within the DNA sample. The methods are used for the CC analysed in the set within the DNA sample. The methods are used for the CC detection of short tandem repeats as genetic markers for the development CC genes, and the simplification and precision of DNA typing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 8; Page 77; 122pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genetic markers
diseases genes a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Simultaneous amplification of short tandem repeats - used to provide genetic markers for linkage maps, for identifying and characterising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Schumm
                WPI; 2000-400106/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA47246 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 23
                                              Schumm JW, Sprecher CJ;
                                                                                                                                                                                                                                                                                                           Primer; short tandem repeat; STR; multiplex amplification reaction; Combined DNA Index System; CODIS; paternity test; breeding; forensi
                                                                                                                                                                                                                                                                                                                                                             Primer 1 for human genomic DNA polymorphic STR locus D22S683
                                                                                                                                                                                                                                                                                                                                                                                                   12-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA47246;
                                                                                                                     25-NOV-1998;
                                                                                                                                                   24-NOV-1999;
                                                                                                                                                                                      02-JUN-2000
                                                                                                                                                                                                                        WO200031306-A2
                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                           profile; D22S683; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1997-526472/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23 ATCTTGGCTCAATGCAACCT 4
                                                                                  PROMEGA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATCTTGGCTCACTGCAACCT 686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genes and for DNA typing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Micka KA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 7 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                     98US-00199542
                                                                                                                                                       99WO-US027876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rabbach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 18.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                               forensic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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New method for analyzing e.g. human tissue DNA samples comprises coamplification of at least 13 short tandem repeat loci, useful in e.g. determining the parentage of a child.

Claim 9; Page 78; 90pp; English.

AAA47201-307 are oligonuclectide primers used to amplify human genomic CC DNA short tandem repeat (STR) loci. The claimed method comprises CC simultaneous determination of the alleles present in a set of loci from CC one or more DNA samples. In particular, at least thirteen loci of genomic CC DNA are amplified in a single multiplex reaction. At least one of the CC loci is preferably a STR locus with a repeat unit of five to seven bases CC or base pairs in length. Preferred loci are thirteen human STR loci CC chosen by the United States Federal Bureau of Investigation as core loci CC chosen by the United States Federal Bureau of Investigation as core loci CC chosen by the United States Federal Bureau of Investigation as core loci CC p381538, HOMTHO1, D21811, D18851, HOMWWFR31, D881179, HOMTPOX, HUMFIERA, CC D588189, D138317, D78820, D168539 and HUMCSB190. Some sets of loci compaining at least one DNA sample; (b) selecting a set of CC minimal incidence of artifacts, e.g. due to repeat slippage. The method CC comprises; (a) obtaining at least one DNA sample; (b) selecting a set of CC continual incidence of artifacts, e.g. due to repeat slippage. The method CC comprises; (a) obtaining at least one DNA sample; (b) selecting a set of CC continual incidence of artifacts, e.g. due to repeat slippage. The method CC comprises; (a) obtaining at least one DNA sample; (b) selecting a set of CC continual continual reaction, the product of the reaction comprising a mixture of amplified alleles from each of the co-amplified loci in the set in a cc continual continua samples found at a crime scene

Sequence 23 BP; 7 A; 4 C; 7 G; 5 T; 0 U; 0 Other;

Query Match Best Local S Matches 19 667 ATCTTGGCTCACTGCAACCT 686 l Similarity 19; Conserv Conservative 1.9%; 0, Score 18.4; Pred. No. 1. Mismatches 1.4e+03DB 1; Length 23; Indels <u>,,</u> Gaps

0

RESULT 907 AAQ25869/c ID AAQ258 AAQ25869 standard; DNA; 19 BP

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23

ATCTTGGCTCAATGCAACCT 4

AAQ25869;

25-MAR-2003 04-JAN-1993 (revised)
(first entry)

3'. Alu primer.

PCR; sequence conservation; DNA synthesis; amplification;

Synthetic.

WO9210566-A1

25-JUN-1992

21-NOV-1991; 91WO-US008739

L3-DEC-1990; 90US-00627945

(TEXA ) UNIV TEXAS SYSTEM

Siciliano

3

Liu

WPI; 1992-234623/28

Chromosome-specific DNA probes free of species-specific repeat for identification and banding of human chromosomes. DNA - used

Claim

65;

Page

63;

73pp; English

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RESULT 908
AAQ25868
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC The sequences given in AAQ25868-9 are nucleotide primers which are CC characterised by binding to a 5′ and a 3′ Alu terminus, respectively. CC These Alu primers were based on a current revision of consensus sequence CC of Alu repeats. This revision is based on nucleotide sequences of 50 CC different, cloned and sequenced human Alu segments. Two regions on the CC sequence showed a high degree of conservation and these were used as CC candidate regions for the primer locations. In order to minimize the CC incorporation of Alu sequence itself in the inter-Alu-PCR, the 5′ primer CC was designed to recognise a specific region and to direct DNA synthesis CC bound. The converse is true for the 3′ primer. Amplification using these CC two primers yields products ranging from a few hundred to several CC thousand base pairs. The primer design maximizes both the number of Alu Segments recruited and the number of inter-Alu unique sequences CC amplified. (Updated on 25-MAR-2003 to correct PN field.)
       X5XFFX8X8X8X8X8X8X8X8X8X8X8X8X8X833333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 17
The sequences given in AAQ25868-9 are nucleotide primers which are characterised by binding to a 5' and a 3' Alu terminus, respectively. These Alu primers were based on a current revision of consensus sequence of Alu repasts. This revision is based on nucleotide sequences of 50 different, cloned and sequenced human Alu segments. Two regions on the sequence showed a high degree of conservation and these were used as candidate regions for the primer locations. In order to minimize the incorporation of Alu sequence itself in the inter-Alu-PCR, the 5' primer was designed to recognise a specific region and to direct DNA synthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 3 A; 8 C; 3 G; 3 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5' Alu primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2003
04-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ25868 standard; DNA; 19
                                                                                                                                                                                                             Chromosome-specific DNA probes free of species-specific repeat DNA for identification and banding of human chromosomes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR; sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ25868;
                                                                                                                                                                            Claim 64; Page 63; 73pp; English.
                                                                                                                                                                                                                                                                                                  Siciliano MJ,
                                                                                                                                                                                                                                                                                                                                                                     13-DEC-1990;
                                                                                                                                                                                                                                                                                                                                                                                                         21-NOV-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                            25-JUN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9210566-A1
                                                                                                                                                                                                                                                                                                                                    (TEXA ) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                645 CAGGCTGGAGTGCAGTGGC 663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CAGGCTGGAGTGCARTGGY 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                      TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                conservation; DNA synthesis; amplification;
                                                                                                                                                                                                                                                                                                                                                                                                        91WO-US008739
                                                                                                                                                                                                                                                                                                                                                                       9008-00627945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18.2;
No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                 used
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RESULT 910 AAQ48683/c ID AAQ486 XX AC AAQ486

AAQ48683 standard; cDNA; 19

BP.

AAQ48683

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                              Matches
                                                        Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               off the 5' end and away from the middle of the Alu segment to which it is bound. The converse is true for the 3' primer. Amplification using these two primers yields products ranging from a few hundred to several thousand base pairs. The primer design maximizes both the number of Alu segments recruited and the number of inter-Alu unique seguences amplified. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                           Brook
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 5 A; 3 C; 6 G; 3 T; 0 U;
                                                                                                                 The sequence is that of a PCR primer Alu-1 which specifically recognises human consensus sequences located at the 5' and 3' ends of Alu segments. It was used with 2F5 template to amplify human unique sequences. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                           Sequence 19 BP; 5 A; 3 C; 6 G; 3 T; 0 U; 2 Other;
                                                                                                                                                                                                  DNA sequence of identify CHR 19
                                                                                                                                                                                                                                                                                                                                                        02-SEP-1993.
                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                            Abnormality;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human Alu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ48682;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ48682 standard;
                                                                                                                                                                           Example; Page 32; 64pp; English.
                                                                                                                                                                                                                                      WPI; 1993-288410/36
                                                                                                                                                                                                                                                                                                           20-FEB-1992;
                                                                                                                                                                                                                                                                                                                                 19-FEB-1993;
                                                                                                                                                                                                                                                                                                                                                                               WO9317104-A1
                                                                                                                                                                                                                                                                                   (MASI ) MASSACHUSETTS INST TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      868 GGATTACAGGCGTGAGCCA 886
                       868 GGATTACAGGCGTGAGCCA 886
                                                                                                                                                                                                                                                             Ą,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -
                                             1 Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GGATTACAGGYRTGAGCCA
  GGATTACAGGYRTGAGCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                    segment
                                                                                                                                                                                                                                                             Housman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                            polymerase chain reaction; amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
                                                                                                                                                                                                                                                                                                           92US-00839255
                                                                                                                                                                                                                                                                                                                                  93WO-US001545
                                                                                                                                                                                                   myotonic dystrophy gene abnormality and protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                      consensus
                                                                                                                                                                                                                                                             DE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.8%;
89.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
                                                           89.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence PCR primer Alu-1.
                                              2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 18.2;
  19
                                                                    Score 18.2;
                                                           Pred.
                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                           No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.3e+03
                                                         1.3e+03

    used to produce probes
kinase reponsible.

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 Other;
                                                                       DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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                                                                    Length 19;
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                                                 Indels
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                                                 Gaps
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RESULT 911
AAQ85677/c
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Best Local Similarity
Matches 17; Conserv
         01-SEP-1993;
                                                                                                                                    Synthetic.
                                                                                                                                                                                             25-MAR-2003
04-OCT-1995
                                                                                                                                                                                                                        AAQ85677;
                                                                                                                                                                                                                                           AAQ85677 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                       Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                         The sequence is that of a PCR primer Alu-2 which specifically recognises human consensus sequences located at the 5' and 3' ends of Alu segments. It was used with 2F5 template to amplify human unique sequences. (Updates on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                         Example; Page 32; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA sequence of myotonic dystrophy gene - used to produce probes identify CHR 19 abnormality and protein kinase reponsible.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1993-288410/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Brook JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-FEB-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-FEB-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Abnormality;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human Alu segment consensus sequence PCR primer Alu-2.
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25-FEB-1994
                          01-SEP-1994;
                                              09-MAR-1995.
                                                                  WO9506714-A1
                                                                                                      misc_difference
                                                                                                                                                       Wilson's disease; chromosome 13; Alu;
                                                                                                                                                                        PCR primer alu 2 for inter-Alu region of Wilson's disease gene
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                                                                                                                                                                                                                                                                                            19
                                                                                                                                                                                                                                                                                                    CAGGCTGGAGTGCAGTGGC 663
                                                                                                                                                                                                                                                                                           CAGGCTGGAGTGCARTGGY 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Housman DE
                                                                                                                                                                                                                                                                                                                                                                       BP; 3 A; 8 C; 3 G; 3 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polymerase chain reaction; amplification;
                                                                                                                                                                                           (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  92US-00839255.
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         93US-00118441
                           94WO-US009851.
                                                                                                       1. .19
                                                                                    /*tag=
/note=
                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                          1.8%;
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                                                                                    "Std IUPAC codes used"
                                                                                                                                                                                                                                            BP
                                                                                                                                                                                                                                                                                                                                 2;
                                                                                                                                                                                                                                                                                                                                 Score 18.2; D
Pred. No. 1.3e
2; Mismatches
                                                                                                                                                       PCR
                                                                                                                                                                                                                                                                                                                                           .3e+03;
                                                                                                                                                       primer;
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RESULT 912
AAQ85676
ID AAQ856
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          In order to physically map and clone the region of the Wilson's disease (WD) gene, a 4.3kb insert from the WD flanking marker D13S31 (probe pCR13Z4) was used to screen a large insert, CBEH II YAC sublibrary. A higher resolution YAC map was constructed using inner-Alu PCR product from 4 large YAC clones to screen the 1431 colony CEPH I YAC sublibrary. A total of 16 mid-size YACs were identified. The pattern of mid-size YACs detected by each large YAC clone was used to order the smaller YAC clones relative to one another. Inter-Alu PCR "fingerprinting" of YAC clones further assisted the ordering process. The data for this are not given in the publication. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                          Gilliam TC,
                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                         Wilson's disease; chromosome 13; Alu;
                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
04-OCT-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQ85676;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ85676 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated Wilson's disease nucleic acid mol. - also probes, vectors, etc., useful for diagnosis and gene therapy of Wilson's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1995-115430/15.
Example; Page 30; 175pp; English.
                       Isolated Wilson's disease nucleic acid mol. - also probes, vectors, useful for diagnosis and gene therapy of Wilson's disease.
                                                                 WPI; 1995-115430/15.
                                                                                                                    (UYCO )
                                                                                                                                                            01-SEP-1993;
                                                                                                                                                                                     01-SEP-1994;
                                                                                                                                                                                                               09-MAR-1995
                                                                                                                                                                                                                                         WO9506714-A1
                                                                                                                                                                                                                                                                                          misc_difference
                                                                                                                                                                                                                                                                                                                                                                                     PCR primer alu 1 for inter-Alu region of Wilson's disease gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 Similarity
17; Conserv
                                                                                                                    UNIV COLUMBIA NEW YORK. GEN HOSPITAL CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.8%;
llarity 89.5%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tanzi RE;
                                                                                          Tanzi RE;
                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
(first entry)
                                                                                                                                                            93US-00118441.
                                                                                                                                                                                     94WO-US009851
                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                   /note=
                                                                                                                                                                                                                                                                              /*tag=
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"Std IUPAC codes used"
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                             PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                            primer;
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RESULT 913
AAQ76249/c
ID AAQ762
XX AAQ762
XX AAQ762
XX 25-MAR
DT 25-MAR
DT 10-AUG
XX Primex
KW Primex
KW Alu cc
KW Alu cc
KW Chroni
XX Synthe
XX Synthe
PN WO942E
XX Synthe
PN WO942E
XX Synthe
IN WO942E
XX Synthe
CX WPI,
IN COLUMN
PR 01-JUN
XX ON-JUN
XX ON-JUN
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XX ON-JUN
PR 01-JUN
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Best Local
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                        The consensus sequence, from bases 13-31, of the 5' end of a 300 bp Alu segment. The sequence was used to generate a set of primers, designated Alu-1 primers set (AAQ76247). The primers of the set have a reverse complementary sequence to the Alu consensus sequence. Thus priming with the Alu-1 set directs synthesis towards the 5' end (1.e. away from the middle) of the Alu segment. Since the primer set is designed to bind close to the edge of an Alu segment, amplification with these primers will reduce the amount of Alu segment sequence and increase the amount o specific chromosomal DNA present required for probe production. The primer set is useful in the production of chromosomal specific probes esfor the detection of chromosomal breakpoints and rearrangements such as probe to detect chronic myelogenous leukemia characterised by the philadelphia chromosome, arising from a reciprocal translocation t (9;22)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA probe specific for Human chromosome region bcr/abl rearrangement in interphase minlei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primer; PCR; amplification; primer set; probe; Alu sequence; Alu repeat; Alu consensus sequence; chromosome; breakpoint; rearrangement; chronic myelogenous leukemia; Philadelphia chromosome; translocation; ss
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10-AUG-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUN-1994;
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Pred. No. 1.3e-
2; Mismatches
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(q34;q11).

(Updated on 25-MAR:2003

to correct

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RESULT 914
AAQ76247
ID AAQ762
XX AAQ762
XX AAQ762
XX DT 10-AUU
XX Gener:
XX Primes
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KW Alu co
KW Chron
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Best Local Similarity
                                                     Matches
                                                                          Query Match
Best Local
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10-AUG-1995
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                                                                                                                                                                                                               The generic sequence of a primer set designated Alu-1. The primer set was based on bases 13-31 of the 5' end of a 300 bp Alu segment (AAQ76249). The primers of the set have a reverse complementary sequence to the Alu consensus sequence. Thus priming with the Alu-1 set directs synthesis cowards the 5' end (i.e. away from the middle) of the Alu segment. Since the primer set is designed to bind close to the edge of an Alu segment, amplification with these primers will reduce the amount of Alu segment sequence and increase the amount of specific chromosomal DNA present required for probe production. The primer set is useful in the production of chromosomal specific probes e.g for the detection of chromosomal breakpoints and rearrangements such as a probe to detect chronic myelogenous leukemia characterised by the Philadelphia chromosome, arising from a reciprocal translocation t(9;22) (q34;q11). (Updated on 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA probe specific for Human chromosome region 9q34 - allows bcr/abl rearrangement in interphase nuclei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Generic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ76247;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ76247 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Siciliano MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9428178-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Primer;
                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 11; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (TEXA ) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mer; PCR; amplification; primer set; probe; Alu sequence; Alu repeat;
consensus sequence; chromosome; breakpoint; rearrangement;
onic myelogenous leukemia; Philadelphia chromosome; translocation; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             868 GGATTACAGGCGTGAGCCA
     868
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer from Alu-1 primer set.
                                                                            Similarity
     GGATTACAGGCGTGAGCCA 886
                                                                                                                                                  19
                                                                                                                                                                                                   to correct PN field.)
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                                                                                                                                                  B₽;
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(first en
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                                                                                                                                                  5 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.8%;
                                                                                                                                                  3 C; 6 G; 3 T;
                                                                            1.8%;
89.5%;
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pred. No. 1.3e
2; Mismatches
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                                                                                                    Score 18.2;
                                                                                  Pred.
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                                                        Mismatches
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                                                                            .3e+03
                                                                                                                                                       2 Other
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                                                                                                      DB 1;
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RESULT 915
AAV83937
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                               RESULT 916
                                                                                                                                                                                                                                                CC produce yeast artificial chromosome (YAC) probes. The YAC probes are used to isolate the nucleic acid sequences of the invention. The specification concluding a necessary sequences derived from a eukaryotic chromosome, are able, in a compatible cell, of replicating acting acting a extractic chromosomal element and segregating during cell division. The sequences concluding a necessary sequences for use in gene therapy, comprising a replicable, segregating during cell division. The sequences can be used to construct artificial chromosomes for use in gene therapy, comprising a replicable, segregating nucleic acid that confers a specific phenotype on cells. Human artificial chromosomes can propagate in human cells and carry large amounts of DNA (e.g. therapeutic genes), and, being extra-chromosomal, they are not mutagenic. The artificial chromosomes are consuction of proteins and to make diagnostic resgents, end. Despression of cytokines, receptors and growth factors, or to increase the copy number of a gene in a cell. The constructs may also be used for functional and structural analysis of chromosomes
                                                                                                                                                  Best Loc
Matches
                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Choo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yeast artificial chromosome; YAC; probe; eukaryotic chromosome; neocentromere; replication; extra-chromosomal element; segregation; cell division; artificial chromosome; gene therapy; human artificial chromosome; transgenic; PCR primer; ss.
AAX09336 standard;
                                                                                                                                                                                                                    Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated nucleic acid comprising neocentromere sequences from eukaryotic chromosome - used to produce replicable, segregating artificial chromosomes that can carry large amounts of DNA for generations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV83937 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 24; 540pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (AMRA-)
                                                                                                                                                  Local Similarity
les 17; Conserv
                                                                                                                   868 GGATTACAGGCGTGAGCCA 886
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                                                                                   GGATTACAGGYRTGAGCCA 19
                                                                                                                                                                                                                    BP; 5 A; 3 C; 6 G; 3 T; 0 U; 2 Other;
                                                                                                                                                  Conservative
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DNA;
                                                                                                                                                                 89.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cancilla MR
18
86
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                                                                                                                                                                  Score 18.2; DB 1;
Pred. No. 1.3e+03;
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                                                                                                                                                  Mismatches
                                                                                                                                                  <u>,</u>
                                                                                                                                                                                Length 19;
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RESULT 917
AAV74139
ID AAV741
XX
AC AAV741
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DT 12-APR
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DT 12-APR
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DE Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC spherocytosis, von Willebrand's disease, tuberous sclerosis, hereditary CC spracytosy, von Willebrand's disease, tuberous of the allelic polymorphic markers found in the human CC genome (represented in AAXIO269-X12937). These primers can be used in a CC method for determining polymorphic forms in an individual for use in e.g. CC forensics, paternity testing or for phenocypic typing for diseases such CC as agammaglobulinemia, diabetes insipidus, lesch-Nyhan syndrome, muscular CC typtocyphy, Wiskott-Aldrich syndrome, Fabry's disease, familial CC hypercholesterolemia, polycystic kidney disease, familial CC hypercholesterolemia, polycystic kidney disease, hereditary CC spherocytosis, von Willebrand's disease, tuberous sclerosis, hereditary CC spherocytosis, von Willebrand's disease, tuberous sclerosis, hereditary CC syndrome, osteogenesis imperfecta, acute intermittent porphyria, CC sutchmune diseases, inflammation, cancer, diseases of the nervous CC system, infection by pathogenic microorganisms, and characteristics such CC as longevity, appearance (e.g. baldness, obesity), strength, speed, CC endurance, fertility, and susceptibility or receptivity to particular CC drugs or therapeutic treatments. The isolated polymorphic nucleic acid CC segments can also be used to produce medicaments for the treatment or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       detection; phenotypic typing; characteristic; infection; hereditary; autoimmune disease; cancer; inflammation; drug; therapy; medicament; treatment; marker; primer; 88.
   Human FLAME-1 PCR primer Mchx-pr1.
                                                                                                                                                                                                      AAV74139 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-NOV-1997;
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                                                                      12-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 15; Page 73; 310pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               determining polymorphic forms for use in testing or phenotypic typing for disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated nucleic acid segments from the human genome - used for determining polymorphic forms for use in e.g. forensics, paternity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-286974/25
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                                                                                                                                                                                                                                                                                                                                                                                CCTCCCAAGTAGCTGGGA 18
                                                                                                                                                                                                                                                                                                                                                                                                                       CCTCCCAAGTAGCTGGGA 564
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of such diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 18; DB 1;
Pred. No. 1.3e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
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                                                                                                                                                                                                                                                                                    RESULT 918
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                           This is the nucleotide sequence of primer Mchx-pr1, which is based on a 3' sequence of human EST clone 427786. This clone had been isolated during a database screening for sequences having homology to the FADD-like caspases Mch4 (caspase-10) and MchS/MACH/FLICE. Mchx-pr1 was used with primer Mchx-pr3 (see AAV74140) in a primary PCR of cDNA from a Jurkat cell library. A secondary PCR was performed using primers Mchx-pr2 and Mchx-pr4 (see AAV74141-42). The PCR product was cloned into vector pBluescript II KS+ and used to screen the Jurkat cDNA library. The beta-isoform of FLAME-1 was isolated. This probe was used to isolate a full-length cDNA (see AAV74136) encoding human FLAME-1 (see AAV90107), a novel anti-apoptotic protein useful for developing apoptotic and anti-apoptotic agents for treating e.g. HIV infection, Alzheimer's disease and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated FADD-like anti-apoptotic molecules -apoptotic and anti-apoptotic agents for treating, Alzheimer's disease or neoplastic conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FLAME-1; FADD-like apoptotic/anti-apoptotic molecule;
HIV; infection; Alzheimer's disease; cancer; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9852963-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example; Page 26; 68pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-045296/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alnemri
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-NOV-1998.
                                                                                                                                                                         Human FLAME-1
                                                                                                                                                                                                                               AAV74141;
                                                                                                                                                                                                                                                         AAV74141 standard; DNA; 18 BP
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                                                                                                                                                                                                    12-APR-1999
            20-MAY-1998;
                                      26-NOV-1998
                                                                WO9852963-A1
                                                                                                                                                FLAME-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sapiens
                                                                                         sapiens
                                                                                                                                 3-1; FADD-like apoptotic/anti-apoptotic molecule; human; apoptosis;
infection; Alzheimer's disease; cancer; therapy; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                       208
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                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                AGGCTGGTCTCGAACTCC
                                                                                                                                                                                                                                                                                                                               AGGCTGGTCTCGAACTCC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                  1.8%;
llarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JEFFERSON THOMAS
                                                                                                                                                                                                    (first entry)
                                                                                                                                                                        PCR primer Mchx-pr2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97US-00859167.
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            98WO-US010200.
                                                                                                                                                                                                                                                                                                                                                                                                                                       6 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English
                                                                                                                                                                                                                                                                                                                                                       225
                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                 Score 18;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                             Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       used to develop
e.g. HIV infection,
                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human; apoptosis; PCR; primer; ss.
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RESULT 919
AAZ39610/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This is the nucleotide sequence of primer Mchx-pr2, which is based on a 3' sequence of human EST clone 427786. This clone had been isolated during a database screening for sequences having homology to the FADD-like caspases Mch4 (caspase-10) and Mch5/MACH/FILCE. Mchx-pr2 was used with primer Mchx-pr4 (see AAV74142) in a secondary PCR of cDNA from a JURKat cell library. A primary PCR was performed using primers Mchx-pr1 and Mchx-pr3 (see AAV74139-40). The PCR product was cloned into vector pBluescript II KS+ and used to screen the JURKat cDNA library. The beta-isoform of FLAME-1 was isolated. This probe was used to isolate a full-length cDNA (see AAV74136) encoding human FLAME-1 (see AAM79107), a novel anti-apoptotic protein useful for developing apoptotic and anti-apoptotic agents for treating e.g. HIV infection, Alzheimer's disease and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New isolated FADD-like anti-apoptotic molecules -apoptotic and anti-apoptotic agents for treating, Alzheimer's disease or neoplastic conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-045296/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Alnemri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example; Page 26; 68pp;
                                                Antisense modulation of human cREL
                                                                         WPI; 2000-061889/05
                                                                                                                                                                             18-SEP-1998;
                                                                                                                                                                                                                                US6001652-A.
                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                           Human;
                                                                                                                                                                                                                                                                                                                                   Human cREL mRNA inhibiting antisense oligo ISIS #24094
                                                                                                                                                                                                                                                                                                                                                                                       AAZ39610;
                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ39610 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYJE-) UNIV
                                                                                                 Monia BP,
                                                                                                                                                   18-SEP-1998;
                                                                                                                                                                                                      14-DEC-1999.
                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                              28-FEB-2000
                                                                                                                           (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     851 GGCCTCCCAAAGTGCTGG 868
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
18; Conserv
                                                                                                                                                                                                                                                                                                           crel;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGCCTCCCAAAGTGCTGG 18
                                                                                                 Cowsert LM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JEFFERSON THOMAS
                                                                                                                           PHARM INC
                                                                                                                                                                                                                                                                                                           transcriptional
                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                    98US-00156253
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                English
                                                                                                     Baker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 18;
Pred. No.
                                                                                                                                                                                                                                                                                                            activator; antisense compound; therapeutic;
                                                                                                    BF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                 expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Le
1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          to develop
HIV infection,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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Gaps

0

The invention provides antisense compounds targeted to a coding region,

Claim 1; Col

27; 26pp;

English.

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RESULT 920
AAH38730/c
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Best Local S
Matches 18
Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPS. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleix acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (transcriptional activator). The antisense compounds are useful as research agents and diagnostics such as in the elucidation of the function of a particular gene. The antisense compounds can be useful as therapeutic modalities that can be configured to be useful in treatment regimes for treatment of cells, tissues and animals, especially humans. In the prior art, there are no known therapeutic agents which effectively inhibit the synthesis of CREL and additional agents capable of inhibiting CREL function are still required. Sequences AAZ39588-627 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; scute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH38730 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 57; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP specific lower PCR primer SEQ ID 1526.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-290930/30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-OCT-2000; 2000WO-US028436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Single nucleotide polymorphism; SNP; single nucleotide primer extension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ORCH-) ORCHID BIOSCIENCES INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CAAAGTGCTGGGATTACA 405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5'UTR of a nucleic acid molecule encoding human cREL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CAAAGTGCTGGGATTACA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ഗ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  股
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.3e+03;
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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to

Claim 1; Page 59; 83pp; English

New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc acid sample.

in a nucleic

WPI; 2001-290930/30 Picoult-Newburg L,

(ORCH-) L5-OCT-1999;

ORCHID BIOSCIENCES INC

Pohl M;

99US-0160096P

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RESULT 921
AAH38990/c
ID AAH3889
XX AAH38990/c
ID AAH388
XX AAH389
AC AAH388
AC AAH388
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AC AAH388
AC AAH388
AC AAH389
AC AAH388
AC AAH38

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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular trophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
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The invention relates to FADD-like apoptotic/anti-apoptotic proteins (FLAME 1 or 2) and nucleic acid molecules encoding such proteins. FLAME sequences are useful for inhibiting apoptosis and for gene therapy of diseases characterised by apoptosis including HIV infection and Alzheimer's disease. FLAME inhibitors are useful as apoptotic agents and activators are useful as anti-apoptotic agents. FLAME-1 is useful as a substrate for caspase in assays to identify caspase inhibitors. The present sequence is human FLAME-1 specific PCR primer, used in the
                                                                                                                                                                                                                                                                                            Novel FADD-like apoptotic/anti-apoptotic proteins useful for inhibiting apoptosis, treating diseases characterized by apoptosis e.g. HIV infection and Alzheimer's disease, and for identifying modulators of the
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3-1; PCR; primer;
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nerapy; human immunodeficiency virus; HIV infection; apoptosis;
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                                                                                                          The invention relates to FADD-like apoptotic/anti-apoptotic proteins (FLAME 1 or 2) and nucleic acid molecules encoding such proteins. FLAME sequences are useful for inhibiting apoptosis and for gene therapy of diseases characterised by apoptosis including HIV infection and Alzheimer's disease. FLAME inhibitors are useful as apoptotic agents activators are useful as anti-apoptotic agents. FLAME-1 is useful as substrate for caspase in assays to identify caspase inhibitors. The present sequence is human FLAME-1 specific PCR primer, used in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disease;
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                                             Gaps
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of the invention

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ADG32591/c

ID ADG32591;

XC ADG32591;

XX ADG32591;

XX 26-FEB-2004 (first entry)

XX mouse; murine; PCR; ss; ve

KW winflammation; skin disorde

KW vRLS; VRLX; VR4; TRPV7; TR

KW inflammation; skin disorde

KW vRLS; VR1X; VR4; TRPV7; TR

KW inflammation; skin disorde

KW wo2002101045-A2.

XX W20022101045-A2.

XX W20022101045-A2.

XX W20022101045-A2.

XX W200202101045-A2.

XX (NOVS) NOVARTIS AG.

YX (NOVS) NOVARTIS AG.

YX W21; 2003-156962/15.

XX W22; 2002-150032
                                                                                                                                                     RESULT 925
ADH59598
   HXXXX
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                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 18
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22-JAN-2002; 2002US-0351238P.
29-JAN-2002; 2002US-0352914P.
12-FEB-2002; 2002US-0357161P.
15-MAY-2002; 2002US-0381086P.
16-MAY-2002; 2002US-0381739P.
   25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acids and encoded proteins thereof. Specifically, it refers to certain members of the VR family that are involved in pain perception, in particular, TRPV3 (previously knows KNLS, VRLX, VR4 & TRPV7), TRPV4 (previously known as VRL3 & OTRPC4) and TRPM8 (previously known as TRPX). Furthermore, this invention includes trkh+ pain specific genes expressed in the sensory neurons of the dorsal root ganglia. Accordingly, such compositions can be useful for the diagnosis, treatment and prevention of pain, inflammation, skin disorders and cancer, and so exhibit analgesic, antiinflammatory, dermatological and cytostatic activities. This oligonucleotide sequence is a PCR primer used to amplify the murine TRPV3
                                                                 ADH59598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated TRPV3, TRPV4 or TRPM8 vanilloid receptor nucleic acid molecule and polypeptides, useful for the diagnosis and treatment of disorders such as pain, inflammation, skin diseases and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Murine TRPV transcript PCR primer SeqID 46.
                                                                                                                           ADH59598 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                dermatological; cytostatic; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to novel vanilloid receptor (VR) related nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; SEQ ID NO 46; 197pp; English.
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VRLS; VRLX; VR4; TRPV7; TRPV4; VRL3; OTRPC4; TRPM8; TRPX; trkA+;
inflammation; skin disorder; cancer; analgesic; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                               638 TGTCACCCAGGCTGGAGT 655
                                                                                                                                                                                                                                                                                    18
                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                    TGTCACCCAGGCTGGAGT 1
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(first entry)
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Pred. No. 1.3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 18;
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Query Match Best Local S Matches 18

Similarity

100.0%;

Score 18; Pred. No.

DB 1; 1.3e+03;

Length 18;

<u>.</u>

Gaps

0

<u>.</u>;

Sequence 18

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3 C; 8

G; 4 T; 0 U; 0

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Comprises contacting the sample with the mixture of probes (preferably comprises contacting the sample with the mixture of probes (preferably comprises contacting the sample with the one or more detectable nucleic acid probes, and determining the target genomic nucleic acid of the sample by determining the hybridization of the one or comore detectable nucleic acid probes to the target genomic nucleic acid of the sample. The genomic nucleic acid is contained in a fixed tissue or a cell, and the sample is metaphase spreads, interphase nucleic or nucleic found in paraffin embedded tissue material or frozen tissue sections. The probe is also useful in comparing a sample of genomic nucleic acid with that of a control sample using a genomic nucleic acid and control genomic nucleic acid and the array or both the sample and control genomic nucleic acid and contacting the array with treated mixture of sample and control genomic nucleic acid and contacting the array with treated mixture of sample and control genomic nucleic acid under suitable hybridization conditions, and comparing the intensities of the signals from the differential labels of the array to that caused by hybridization of the probes to genomic nucleic acid, thus determining one or more variations in copy numbers of sequences in the control. The hybridization of the genomic conditions, and comparing the probes to genomic nucleic acid, thus determining the promote nucleic acid hybrid. The sample as compared with the relative copy numbers of substantially corrected to sample as compared with the relative copy numbers of substantially corrected genomic nucleic acid hybrid. The hybridization of the genomic nucleic acid hybrid. The sample acid are labelled with detectable moiety such that hybridization of the genomic nucleic acid to be tested and the reference of nucleic acid and the reference of nucleic acid that is prepared from the one or more genomic nucleic 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Non-nucleotide probe for suppressing binding of detectable nucleic acid probes to undesired sequences, has aggregate nucleobase sequence homologous to randomly distributed repeat sequence of genomic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents a non-nucleotide probe. The probe is useful for suppressing the binding of one or more detectable nucleic accupables, that are greater than 100 base pairs and that have been derived from genomic nucleic acid, to one or more undesired sequences in an ass.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 10; SEQ ID NO 4; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              non-nucleotide probe;
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represents a non-nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for determining target
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DAKOCYTOMATION DENMARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hyldig-Nielsen JJ, Williams
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     genomic nucleic acid of a sample.
probe
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          of the invention.
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RESULT 926 ADH59610/c ID ADH596 ADH59610 standard; DNA; 18 ADH59610; В₽

Non-nucleotide probe of the invention #14 25-MAR-2004 (first entry)

probe. non-nucleotide probe; Bacterial Artificial Chromosome clone; BAC; ss;

Synthetic

WO2003027328-A2

24-SEP-2002; 2002WO-US030573

24-SEP-2001; 2001US-0324499P (BOST-) BOSTON PROBES

(DAKO-) DAKOCYTOMATION DENMARK AS

, VV, Hyldig-Nielsen JJ, Williams

WPI; 2003-421160/39

Non-nucleotide probe for suppressing binding of detectable nucleic acid probes to undesired sequences, has aggregate nucleobase sequence homologous to randomly distributed repeat sequence of genomic nucleic

Claim 10; SEQ ID NO 16; 103pp; English

CC more detectable nucleic acid probes to the target genomic nucleic acid of the sample by determining the hybridization of the one or common the control sample. The genomic nucleic acid is contained in a fixed tissue or a ccell, and the sample is metaphase spreads, interphase nucleic or nucleic found in paraffin embedded tissue material or frozen tissue sections. The ccell of a control sample using a sample of genomic nucleic acid with that of a control sample using a genomic nucleic acid reference array. The method comprises treating a sample of genomic nucleic acid and control genomic nucleic acid and the array with the mixture of the probe under suitable hybridization conditions, contacting the array with treated mixture of sample and control genomic nucleic acid and the array to contacting the signals from the differential labels of the array to that caused by hybridization of the probes to genomic nucleic acid, thus determining one or more variations in copy numbers of sequences in the control. The hybridization of the genomic nucleic acid, thus compared with the relative copy numbers of substantially contacting the control sequences in the control. The hybridization of the genomic conditions, and comparing the control sequences in the control. The hybridization of the genomic of the genomic of the genomic array is determined using an intercalating dye or a detectable antibody, or its fragment, that is specific for a nucleic acid, thus detectable antibody. The sample of genomic array is determined by determining the presence, absence, amount or location of the detectable moiety such that hybridization of the detectable useful for suppressing the binding of one or more detectable nucleic probes, that are greater than 100 base pairs and that have been derived from genomic nucleic acid, to one or more undesired sequences in an after determining target genomic nucleic acid of a sample. The method comprises contacting the sample with the mixture of probes (preferable comprising 5-50 probes), contacting the sample with the one or more detectable nucleic acid probes, and determining the target genomic The present sequence represents a non-nucleotide probe. The probe is probes (preferably the one or more derived

Best Matches Query Match

Local

Similarity

100.0%;

1.8%;

Score 18; ; Pred. No.

DB 1; Length 18;

1.3e+03;

Indels

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Gaps

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18;

Conservative

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Mismatches ŏ.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                        The present invention relates to a neural thread protein (NTP) peptide referred to as cell death peptide. Thought to be cytostatic, antibacterial, immunosuppressive and antiinflammatory. It is useful for treating a condition in a patient requiring removal or destruction of cells, for treating a condition such as benign or malignant tumor, inflammatory disease, autoimmune disease and infectious disease. The peptide useful for treatment is derived from the amino acid sequence for a pancreatic thread protein. The peptide is conjugated, linked or bound to a molecule chosen from antibody or its fragment, antibody-like binding molecule, where the molecule has a higher affinity for binding to a tumor or other target than binding to other cells. Treatment using NTP peptides can remove benign tumors with less risk and fewer of the undesirable side effects of surgery. The present sequence is an NTP encoding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     arrays. The genomic array comprises nucleic acid that is prepared from Bacterial Artificial Chromosome (BAC) clones. The present sequence
                                                                                                                                                                                                                                                                                 Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatos atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-JUL-2001; 2001US-0306150P.
19-JUL-2001; 2001US-0306161P.
16-NOV-2001; 2001US-0331477P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 4 A; 8 C; 3 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    neural thread protein; NTP; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cytostatic; Antibacterial; Immunosuppressive; Antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-AUG-2003
Sequence 18
                                                                                                                                                                                                                                                     Disclosure; Page 19; 77pp; English.
                                                                                                                                                                                                                                                                                                                                               WPI; 2003-247999/24.
P-PSDB; ABR63264.
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 BP; 3
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 0 C; 1 G; 14 T; 0
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Pred. No.
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   U; 0 Other;
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                                                                                                                                                                                                                                                                                                      dermatosis,
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                                          RESULT 929
AAH37310/c
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                                                                                                                                                                                                                   RESULT 928
ACC84468
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 8XB
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                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                           Matches 18;
                                                                                                                                                                                                                                                         The present invention relates to a neural thread protein (NTP) peptide referred to as cell death peptide. Thought to be cytostatic, antibacterial, immunosuppressive and antiinflammatory. It is useful for treating a condition in a patient requiring removal or destruction of cells, for treating a condition such as benign or malignant tumor, inflammatory disease, autoimmune disease and infectious disease. The peptide useful for treatment is derived from the amino acid sequence for a pancreatic thread protein. The peptide is conjugated, linked or bound to a molecule chosen from antibody or its fragment, antibody-like binding molecule, where the molecule has a higher affinity for binding to a tumor or other target than binding to other cells. Treatment using NTP peptides can remove benign tumors with less risk and fewer of the undesirable side
AAH37310
                           AAH37310 standard;
                                                                                                                                                                                                                   Sequence 18
                                                                                                                                                                                                                                             effects of surgery. The present sequence is an NTP encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatosis, atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Averback PA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JUL-2001; 2001US-0306150P
19-JUL-2001; 2001US-0306161P
16-NOV-2001; 2001US-0331477P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       neural thread protein; NTP; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cytostatic; Antibacterial; Immunosuppressive; Antinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-AUG-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 18; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-JUL-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2003-247999/24.
DB; ABR63263.
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                                                                                                  TTTAATTTTTTTTTTTTTT
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                                                                                                                                                                                                                   BP; 2 A; 0 C; 2 G; 14 T; 0 U; 0 Other;
                                                                                                                                                           Conservative
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                           DNA; 19
                                                                                                                                                                      1.8%;
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                                                                                                                                                                       Score 18;
Pred. No.
                                                                                                                                                           Mismatches
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                                                                                                                                                                                        DB 1;
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                                                                                                                                                           0,
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AAH91092/c ID AAH910 XX RESULT 930

AAH91092

standard; DNA; 19

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19

645 CAGGCTGGAGTGCAGTGG

662 ٥,

Matches Best Query Match

Local

l Similarity 18; Conserv

Conservative

1.8%;

Score 18; Pred. No.

DB 1; 1.3e+03

Length 19; Indels

Mismatches

0;

Gaps

0

Sequence

19 BP; 4

A; 9 C; 3 G; 3 T; 0 U; 0 Other;

microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a PCR primer specif:

represents a PCR primer specific

containing DNA

sequence

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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking CC sites of single nucleotide polymorphisms SNPs. The present invention CC includes kits for determining the presence or absence of a SNP, using the CC oligonucleotides of the invention. The PCR primers are used to amplify a CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The CC identity of a SNP and for genotyping nucleic acid sample by coligonucleotides are useful for determining the presence, absence or CC identity of a SNP and for genotyping nucleic acid sample by cassess by association analysis the genotype of an individual or group of casses by association analysis the genotype of an individual or group of casses by one or more SNPs. Phenotypic trait suspected of being CC caused by one or more SNPs. Phenotypic traits include diseases e.g. CC agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular CC diseases imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial CC diseases, including, rheumatoid arthritis, multiple sclerosis, pathogenic conformation. The method is also useful in forencic investigations and conformation and conformation in also useful in forencic investigations and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Single nucleotide polymorphism; SNP; single nucleotide primer extension;
SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer;
Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia;
polycystic kidney disease; osteogenesis imperfecta; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 acid sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-290930/30.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    50; 83pp; English
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RESULT 931
AAH91352/c
ID AAH913
XX AAH913
XX O9-OCT
XX Human
XX Homo SX Homo 8
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human inflammatory bowel disease associated polymorphic site #167
                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a method for detecting the presence polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to the presence of genetic polymorphisms associated with inflammatory because and correlating their occurrence with disease states. They method the presence of genetic polymorphisms associated with inflammatory because and correlating their occurrence with disease states.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Testing for the bowel disease,
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                                       Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis; single nucleotide polymorphism; SNP; chromosome 19p13; paternity test;
                                                                                                                                                        AAH91352 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                    polymorphic
                                                                                                                                                                                                                                                                                                                                                                   testing,
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10-APR-2000; 2000US-0196046P.
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  Homo sapiens
                                                                             Human inflammatory bowel disease associated polymorphic site
                                                                                                      09-OCT-2001
                            chromosome 5q31-33; forensic test; gene therapy; ds
                                                                                                                                                                                                                                                                                                                                                    in this way for phenotypic correlations, forensics, paternity ing, medicine and genetic analysis. The present sequence is a norphic site described in the exemplification of the invention
                                                                                                                                                                                                                                                769
                                                                                                                                                                                                                      19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inflammatory bowel disease; Crohn's disease; ulcerative colitis; nucleotide polymorphism; SNP; chromosome 19p13; paternity test; some 5q31-33; forensic test; gene therapy; ds.
                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for the presence of polymorphisms associated with inflammatory isease, using a hybridization assay.
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/note= "SNP,
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                                                                                                                                                                                                                                                                                   1.8%;
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Pred. No. 1.3e+03;
D; Mismatches 1
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Best Local
13-SEP-1999;
16-SEP-1999;
20-SEP-1999;
13-OCT-1999;
12-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                              disease and correlating their occurrence with disease states. They used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Testing for the prese
bowel disease, using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Key
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                                                                                                                                                                                                                                                                                                                                                                                                                            polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to the presence of genetic polymorphisms associated with inflammatory disease and correlating their occurrence with disease states. They
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                                                                                                                                                                                                                                                                                                                                                                          Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a method for detecting the presence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 56; 463pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Daly M,
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10-APR-2000;
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                                                                                                     WO200119856-A2
                                                                                                                        Homo
                                                                                                                                                      Human secreted
                                                                                                                                                                          Forward
                                                                                                                                                                                               04-JUL-2001
                                                                                                                                                                                                                   AAS01233;
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                                                            13-SEP-2000;
                                                                                 22-MAR-2001.
                                                                                                                                            PCR primer;
                                                                                                                        sapiens
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                                                                                                                                                                                                                                                                                                                                     18;
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                                                                                                                                                                          PCR
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                                                                                                                                                                                                                                                                                                                                              Similarity
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                                                                                                                                             88
                                                                                                                                                                                                                                                                                                                                                                          BP; 7 A; 4 C; 3 G; 4 T; 0 U; 1 Other;
                                                                                                                                                                          primer,
                                                                                                                                                                                                                                                                                                                                     Conservative
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2000US-0196046P.
                                                              2000WO-US025106
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           99US-0153629P.
99US-0154520P.
99US-0154762P.
99US-0159231P.
                                                                                                                                                       protein; therapeutic; diagnostic; human; cancer;
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                                                                                                                                                                                                                                                                                                                                              94.7%;
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2000US-00659634

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RESULT 933
ACAS6212/C
ID ACAS62
XX ACAS62
AC ACAS62
XX Human
XX Human
XX Human
XX Homo 8
XX Chromo
KW Chromo
KW D4S424
XX US2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The sequence represents the Forward PCR primer, used in expression analysis of human secreted protein, POLYS. POLYX nucleic acids, polypeptides and antibodies to POLYX can be used for treating or preventing a POLYX associated disorder in a subject, preferably a human. These can be used in the manufacture of a medicament for treating a syndrome associated with a human disease selected from a POLYX associated disorder, where the therapeutic is a POLYX polypeptide, a POLYX nucleotide or a POLYX antibody. They may also be used to screen for a mucleotide or a POLYX antibody. They may also be used to screen for a
                                             Determining a genotype associated with increased or decreased resistance to familial bipolar affective disorder in a family comprises determining the genotype of e.g., chromosomal regions D4S402 and D4S424.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; genotype determination; familial bipolar affective disorder; chromosomal region linked; locus associated with resistance; D4S402 D4S424; D4S431; D4S404; D1IS394; D1IS39; chromosome marker; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19
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                                                                                                                                                                                                            Ginns EI,
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20-OCT-1997;
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(EGEL/) EGELAND J A.
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Disclosure; Page 11; 79pp; English

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a method of determining a genotype cassociated with increased or decreased resistance to familial bipolar caffective disorder. The method comprises determining the genotype with at cleast one marker of at least one chromosomal region linked to a locus corromosomal regions are included of and localised between D49402 and C4 chromosomal regions are included of and localised between D49402 and C4 chromosomal regions are included of and localised between D49402 and C5 chromosomal regions or D18394 and D1829. The invention also consider that for determining a genotype associated with increased or discloses a kit for determining a genotype associated with increased or correased resistance to familial bipolar affective disorder, where the ckit comprises markers for two or more of the chromosomal regions cited. The method and kit are useful for determining a genotype associated with increased or decreased resistance to familial bipolar affective disorder cin a family affected by bipolar affective disorder, for determining the contribution of these chromosomal regions to bipolar affective disorder in a affective family member, and for assessing an increased or correspond risk of developing bipolar illness for a tested individual from correspond invention.
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                                             The invention relates to a human protein with cancer suppressing function. Also included are claims for: polymucleotides encoding the polypeptide, the recombinant process of producing the polypeptide in treating various diseases, such as cancer, the agonist resisting the polypeptide and its treatment effect and application of the polymucleotides encoding the human protein with cancer suppressing function. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human cancer suppressing protein associated PCR primer
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The invention relates to human POLYX polypeptides and the polynucleotides concoding them. The invention also relates to an antibody that cimmunospecifically binds to a POLYX polypeptide, a method of determining the presence or amount of a POLYX polypucleotide in a sample involving contacting the presence or amount of the probe bound to the polynucleotide and comethod of identifying an agent that modulates the expression or activity of a POLYX polypeptide involving providing a cell expression or activity complete involving providing a cell expression or activity of a POLYX polypeptide involving providing a cell expression where an agent modulates expression or activity of the polypeptide where an agent modulates expression or activity of the polypeptide where an activation, and a method of modulating the activity of a polypeptide compound that binds to the polypeptide in an amount sufficient to compound that binds to the polypeptide in an amount sufficient to compound that binds to the polypeptide in an amount sufficient to compound that binds to the polypeptide in an amount sufficient to compound that binds to the polypeptide in an amount sufficient to compound that binds to the polypeptide in an amount sufficient to compound that binds to the polypeptide in an amount sufficient to compound the activity. The POLYX polypeptides are useful for contacting the presence of or predisposition to a disease associated with altered levels of POLYX DNA or protein in a first mammalian subject, involving measuring the level of expression of DNA or the amount of protein in a sample from the first mammalian subject and comparing the
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                                                                                                                                                                                                                                                                                                                                                                                                        Herrmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 18;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                        Į,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                        Liu
                                                                                                                                                                                                                                                                                                                                                                                                        ×
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                       Yang
                                                                                                                                                                                                                                                                                                                                                                                                        3
                                                                                                                                                                                                                                                                                                                                                                                                        Boldog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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RESULT 936
ADM32249/c
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                                                                                                                                                                Query Match
Best Local S
                                                                                                                                                     Matches
                                                                                                                                                                                                                                                         amount of DNA or protein in a sample from a second mammalian subject known not to have or not be predisposed to the disease, where an alteration in the expression level of DNA or protein in the first subject as compared to the control sample indicates the presence of a predisposition to the disease. The sequences of the invention are useful for treating or preventing a POLYX-associated disorder which involves administering POLYX DNA. A therapeutic such as a POLYX DNA, protein or antibody is useful in the manufacture of a medicament for treating a syndrome associated with a human disease. This sequence represents a PCR primer used to amplify a human POLYX polynucleotide of the invention.
                                                                                                                                                                                                                          Sequence 19 BP; 4 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                644 CCAGGCTGGAGTGCAGTG
                                                                                                                                                     18;
                                                                           N
                                                                                                                                                                        Similarity
                                                                             CCAGGCTGGAGTGCAGTG
                                                                                                                                                   Conservative
                                                                                                                                                                      100.0%;
                                                                                                                                                                                           1.8%;
                                                                             19
                                                                                                                661
                                                                                                                                                     <u>,</u>
                                                                                                                                                                        Score 18;
Pred. No.
                                                                                                                                                       Mismatches
                                                                                                                                                                                           DB 1;
                                                                                                                                                                          1.3e+03;
                                                                                                                                                                                           Length 19;
                                                                                                                                                         Indels
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                                                                                                                                                          Gaps
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ADM32249
                                                                                                                                                         19-FEB-2004.
                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                single nucleotide polymorphism; ss; primer.
                                                                                                                                                                                                     human interleukin-18; IL-18; adult onset still disease; gene
                                                                                                                                                                                                                Human interleukin-18 gene polymorphism related primer,
                                                                                                                                                                                                                            20-MAY-2004
                                                                                                                                                                                                                                         ADM32249;
                                                                                                                                  22-JUL-2002; 2002JP-00212550
                                                                                                                                              22-JUL-2002; 2002JP-00212550
                                                                                                                                                                    JP2004049136-A
                                                                                                                 (HYUB-)
                                                                                                                      (SUGI/)
                                                                                                                SUGIURA S.
HYUBITTO GENOMICS
                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                    DNA;
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detecting Detecting gene polymorphism in interleukin-18 adult onset still disease. gene of human, useful

WPI; 2004-174121/17

Claim 6; SEQ ID NO 6; 61pp; Japanese.

The invention relates to a novel method for detecting a gene polymorphism in a human interleukin (II)-18 gene. The method involves detecting a 9 base insertion between -6311 position and -6310 position, a polymorphism at positions -5890, -5316, -4762, -4675, -3268, -689 and -640 of a polymuclectide which consists of a fully defined sequence of 6640 base pairs as given in the specification, where in the 6640bp polymuclectide, the position 6575 is set to +1 from which numbering is performed. The method is useful for detecting gene polymorphism in II-18 gene of human and for detecting adult onset still disease. This polymuclectide sequence represents a primer of the human interleukin-18 gene of the invention.

Sequence 19 BP; 4 A; 3 C; 7 G; 5 T; 0 U; 0 Other;

Best Loc Matches Query Match Local 18; Similarity Conservative ( 1.8%; Score 18; 100.0%; Pred. No o ; Mismatches ŏ. 1.3e+03; DB 1; Length 19; Indels <u>.</u> Gaps 0

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RESULT 937
ADP09291;
XX
ADP09291;
XX
AC
ADP09291;
XX
DT
26-AUG-2004 (first of the context of the
RESULT 938
AAH38402
ID AAH384
XX
AAH384
XX
DT 14-AUG
XX
DT 14-AUG
XX
DT SNP sp
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                                                                                                                                                                                                                                                                                                                                                                              片
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 18
   SNP specific lower PCR primer SEQ ID 1198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel method for identifying a subject at risk of breast cancer which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a risk of breast cancer, as well as therapeutic and prophylactic treatments that specifically target breast cancer, such as gene therapy. The current sequence is that of an Extend primer of the invention which was used to genotype single nucleotide polymorphisms within human chromograpin B (CHGB; secretogranin property in the contraction of the contraction which was used to genotype single nucleotide polymorphisms within human chromograpin B (CHGB; secretogranin contraction).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying a subject at risk of breast cancer by detecting the presence or absence of one or more nucleotide polymorphic variations, useful for diagnosing, preventing and/or treating breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          breast cancer; cytostatic; gene therapy; human; chromogranin B; CHC secretogranin 1; SCG1; chromosome 20pter-p12; ss; PCR; primer; SNP; single nucleotide polymorphism.
                                                                     14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-NOV-2002; 2002US-0429136P
24-JUL-2003; 2003US-0490234P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 3 A; 9 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 5; Page 103; 286pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-NOV-2003; 2003WO-US037966
                                                                                                                                        AAH38402;
                                                                                                                                                                                                          AAH38402 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SEQU-) SEQUENOM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2004-441082/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               966
                                                                                                                                                                                                                                                                                                                                                                                                                                             646 AGGCTGGAGTGCAGTGGC 663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18
                                                                                                                                                                                                                                                                                                                                                                              19 AGGCTGGAGTGCAGTGGC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AATCTCGGCTCACTGCAA 983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nelson MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  which is located at chromosomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.8%;
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                                                                                                                                                                                                          ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 18; DB 1;
Pred. No. 1.3e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  position 20pter-p12
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RESULT 939
AAH20695/c
ID AAH206
XX
AC AAH206
XX
AC AAH206
XX
DT 13-AUG
XX

AAH20695;

AAH20695

standard;

DNA;

20 ВP

13-AUG-2001

(first entry)

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1038 GATTACGGGCACCTGCCACC 1057

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GATTACAGGCAMCTGCCACC 20

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Matches
                                                   Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia, diabettes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                                                                                                                                                                                                                                      Sequence 20 BP; 5 A; 7 C; 4 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                   diseases, including, rheumatoid arthritis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a PCR primer specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 56; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-290930/30.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-OCT-2000; 2000WO-US028436.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-APR-2001
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                                                                                                                                                                                                                                                                                                                                                                human
18;
                                                                                                                                                                                                                                                                                                                                                                      SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-0160096P.
                                                                                                                                                                                                                                                                                                                                                                      containing DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pohl M
                                                         1.8%;
                                                                Score 18; DB 1;
Pred. No. 1.4e+03;
      Mismatches
                                                                                                                                DB 1;
                                                                                                                         Length 20;
      0
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Gaps
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Human

telomeric

repeat binding

factor 2

oligonucleotide

111423

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RESULT 940
ABK70676/c
ID ABK706
XX
AC ABK706
XX
AC ABK706
XX
DT 15-JUL
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                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel antisense compound (I) 8-30 nucleobases in length targeted to a polynucleotide encoding human telomeric repeat binding factor 2 (II) which specifically hybridizes with, and inhibits the expression of (II). (I) is useful for treating a human having a disease or condition associated with (II) such as premature aging or a hyperproliferative disorder especially cancer, by inhibiting the expression of (II) in human cells or tissues. (I) is useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. The products of the invention have cytostatic activity. This sequence represents an antisense oligonucleotide used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense; phosphorothioate; human; telomeric repeat binding factor 2; inhibitor; premature aging; hyperproliferative disorder; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antisense compounds targeted to nucleic acid encoding telomeric repeat binding factor 2 useful for treating conditions such as premature agin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                           Sequence
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                       15-JUL-2002
                                                                   ABK70676;
                                                                                                          ABK70676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diseases such as cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sapiens
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                                                                                                                                                                                                                                                                                            l Similarity
18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 81; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                             20 BP; 4
                                                                                                          standard;
                                                                                                                                                                                                                                                   GGCTGGAGTGCAGTGGCG
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                                                                                                                                                                                                                                                                                              Conservative
                          (first
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      RESULT 941
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Human hepatocellular carcinoma
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                                                                                                                                                                                                                                                                 28-MAR-2002
                                                                                                                                                                                                                                                                                                      WO200224948-A2
                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                              Human; hepatocellular carcinoma; HCC; chromosome
                 New nucleic acids useful
                                                        WPI; 2002-383197/41.
                                                                                                                                (INRM )
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                                                                                                                                INST PASTEUR.
INSERM INST NAT SANTE & RECH
                                                                                           Marchio A,
                                                                                           Dejean
for in vitro detection of homozygous deletion a hepatocellular carcinoma cell line.
                                                                                                                                                                                                                                                                                                                                                                                                                     (HCC) homozygous deletion PCR
                                                                                                                                      MEDICALE
                                                                                                                                                                                                                                                                                                                                                                                  8p23; ss; primer; PCR
                                                                                                                                                                                                                                                                                                                                                                                                                       primer #28
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The invention relates to an isolated nucleic acid used for detection of human hepatocellular carcinoma (HCC), through homozygous deletion in human chromosome 8p23. The deletion within the 345 kilobase region flanked by the 370L3SP6 and loci markers. Sequences ABK70649-ABK70700 represent PCR pri Disclosure; Page 14; 32pp; English. loci markers. Sequen detect the deletion chromosome 8p23 of indicative of HCC primers used to is located 315117fg8D in vitro detection

Sequence 20 BP; 3 A; 7 C; 4 G; 6 T; 0 U; 0 Other;

Matches Query Match Best Local ( 729 18; Similarity AGTAGCTGGGACTACAGG 746 Conservative 1.8%; 0; Score 18; Pred. No. Mismatches DB 1; Length 20; 1.4e+03; Indels ٥, Gaps 0

ABZ98008; ABZ98008 standard; DNA; 20 ВP

17-OCT-2003

(first entry)

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18

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; Human RANTES oligonucleotide sequence respiratory therapy;

Homo sapiens

WO200285308-A2

31-OCT-2002

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

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RESULT 942
ABZ92737
T X A X A Z X B X 
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Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions initiation codon,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nyce JW, Li Y, Miller S, Tang
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                                  (EPIG-) EPIGENESIS PHARM INC
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L, Shahabuddin
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initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or in a subject of the region of the prophylactic or the respiratory bronchodilation, increasing levels of ubiquinone or in a subject of the region of the prophylactic or the region 
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Miller S,
                                                                                                       lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                      at ftp.wipo.int/pub/published_pct_sequences
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, Tang L,
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L, Shahabuddin S;
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Query Match Best Local : Matches Local Similarity 373 18;  $\vdash$ CCTGCCTCAGCCTCCCAA 390 ccreccreaeccreccaa 18 Conservative 100.0%; 1.8%; 0; Score 18; Pred. No. Mismatches DB 1; 1.4e+03; Length 20; ٥, Gaps 0

Sequence

20 BP; 4 A; 11 C; 2 G; 3 T; 0 U; 0 Other;

밁 Ś

N58473-derived oligonucleotide 29-JUL-2004 ABD28967; ABD28967 standard; DNA; (first entry) 20 SEQ ID 7979.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antianflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; pulmonary transplantation rejection; ss; primer. beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;

Homo sapiens

WO200285309-A2

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P

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RESULT 944
ABD31039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Composition comprises oligo and is administered to reduce the production composition composition or target polypeptide associated with lung airway or lung composition or cancer and can be anti-sense to the corresponding mRNA.

Composition or cancer and can be anti-sense to the corresponding mRNA.

Composition or cancer and can be anti-sense to the corresponding mRNA.

Composition of the invention has antiallergic, antiinflammatory, antiasthmatic, composition has antiallergic, antiinflammatory, antiasthmatic, composition comprises oligo and is administered to reduce the production of treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, allergies and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, composition rejection, pulmonary infections, bronchotics, pulmonary disease, pulmonary hypoproduction are associated composition rejection, pulmonary infections, bronchitis or cancer.

Composition rejection, pulmonary infections, bronchitis or cancer.

Composition rejection, pulmonary disease, pulmonary pain, cystic fibrosis, allergic rhinitis, pulmonary hypoproduction are pulmonary disease, pulmonary hypoproduction and the readuced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system and the readuced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system and the readuced adenosine to the store triangle environment and thereby. To
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Best Local
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Miller S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pharmaceutical composition for treating ast oligonucleotide containing less percentage nucleic acids associated with lung airway of the containing terms of the containing airway of the containing as the containing actions as the containing action and the containing action are contained as the containing action as the containing action actions are contained as the containing action actions as the containing action actions are contained as the containing action action actions are contained as the containing action act
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (EPIG-) EPIGENESIS PHARM
                                                                                           surfactant depletion; antiallergic; antiinflammatory; antiasthmatic;
analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;
                                                                                                                                                                                Human; antisense;
                                                                                                                                                          respiratory tract
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Tang
                                                                                                                                                                                                                                                                                                                                                                                                                            standard; DNA;
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L, Shahabuddin
                                                                                                                                                       bronchoconstriction; allergy; hyposecretion; pain; inflammation; adenosine sensitivity; lung; cancer;
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ed effects
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Pred. No. 1.4e+03;
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to it
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RESULT 945 ADH77439

ADH77439 standard; DNA;

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ADH77439

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                                                          Matches
                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic,
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Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies reducing adenosine sensitivity, levels of adenosine (A) or (A)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pharmaceutical composition for treating ast oligonucleotide containing less percentage nucleic acids associated with lung airway of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 15;
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394 GCTGGGATTACAGGCGTG
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                                                                                  Similarity
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Tang
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                                                                                                                                                                                                  unwanted effects
                                                                                                                                                                                                                        brain, heart,
                                                            Conservative
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L, Shahabuddin
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                                                                                  Score 18;
Pred. No.
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                                                            Mismatches
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adenosine, targ
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                                                                                                        Length 20;
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nammal. The
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                                      RESULT 946
                                                                                                                               Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                      The invention describes a compound 8-80 nucleobases in length targeted to, and which specifically hybridizes with a nucleic acid molecule encoding PTPN12 (protein tyrosine phosphatase, non-receptor type 12), and inhibits the expression of PTPN12. The compound, composition and methods are useful for treating a disease or condition associated with PTPN12, such as a hyperproliferative disorder, e.g. colon cancer, or a metabolic disorder. They are also useful in research and disagnostics for modulating the expression of PTPN12. This sequence represents a human protein tyrosine phosphatase, non-receptor type 12 (PTPN12) antisense
                                                                                                                                                                                                                                                                                                                                                                              New antisense oligonucleotides targeted to a núcleic acid encoding protein tyrosine phosphatase, non-receptor type 12 (PTPN12) useful treating a disease associated with PTPN12, e.g. colon cancer.
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             ADJ59873 standard; DNA;
                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-061282/06.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein tyrosine phosphatase, non-receptor type 12; hyperproliferative disorder; colon cancer; metabolic disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cytostatic; PTPN12 Inhibitor;
                                                                                                                                                                                                                                                                                                                                                         Example 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2003232434-A1
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                                                                                                      643 CCCAGGCTGGAGTGCAGT 660
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                                                                                                                                           Similarity
                                                                                                                                                                                    20
                                                                            CCCAGGCTGGAGTGCAGT
                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 80; 117pp; English.
                                                                                                                                                                                 BP; 3
                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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/mod_ba
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        note=
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                                                                                                                                                                                 Α,
                                                                                                                                           1.8%; Score 18; DB 1; 100.0%; Pred. No. 1.4e+0
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e= "OTHER= 2'-O-methoxyethyl (2'-MOE)
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                                                                                                                                                                                 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "OTHER= 2'-O-methoxyethyl (2'-MOE)
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                                                                                                                                Mismatches
                                                                                                                                           1.4e+03;
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                                                                                                                                                       Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleotides
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RESULT 947
ADM15386/c
ID ADM153
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                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(les), asthma, impeded respiration, cystic fibrosis (CR), chronic obstructive pulmonary diseases (CDPD), allergic rhintis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway the present and oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 4 A; 3 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nyce JW, Tang L,
Shahabuddin S, Lu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oligonucleotide associated to RANTES
                                  01-JUL-2004
                                                                    ADM15386;
                                                                                                     ADM15386 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-203534/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JUL-2002; 2002US-0399076P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADJ59873;
                                                                                                                                                                                                                                                                                                                                                                                     obstruction.
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                                                                                                                                                                                                                           394 GCTGGGATTACAGGCGTG 411
                                                                                                                                                                                          N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EPIGENESIS PHARM INC
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                                                                                                                                                                                          GCTGGGATTACAGGCGTG 19
                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                  (first entry)
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H, Cong H;
                                                                                                                                                                                                                                                                                 100.0%;
                                                                                                                                                                                                                                                                                                  1.8%;
                                                                                                       20
                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                Score 18; DB 1;
Pred. No. 1.4e+03
                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                  Length 20;
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                                                                                                                                                                                                                                                               Gaps
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Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1573

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The present sequence represents a chimeric antisense oligonucleotide crargeted to human microsomal prostaglandin E2 synthase (mPGES-1). The CC human mPGES-1 gene is located on chromosome 9, more specifically to CC 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and CC mPGES-1; by the syression; (2) a method of inhibiting the expression of CC mPGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mPGES-1. MPGES-1 chimeric CC antisense or condition associated with mPGES-1. MPGES-1 chimeric CC antisinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulator, cardiant, neuroprotective, contininflammatory, neuroprotective, nootropic, antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's CC condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition of the condition injury, or condition of the condition injury, or condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition injury, or condition of the condition injury, or condition conditions are composition injury, or conditions are conditions and can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's conditions are conditions of the condition injury, or conditions are conditions of the condition injury, or conditions are conditions of the conditions of the conditions of the condition injury, or conditions are conditions of the condition
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gierse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified_base
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16. .20
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/mod_base= OTHER
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'note= "2'-O-methocyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English.
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Sequence

20

BP; 12

A.

3 C; 0 G; 5 T; 0 U; 0 Other;

Local Similarity

100.0%;

Score 18; Pred. No.

1.4e+03 DB 1;

Length 20

. 88

ophthalmic,

immunological,

cardiovascular

o R

neurological

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RESULT 948
ADO45363
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chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-5 receptor, CCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDB4 A, PDB4 B, PDB4 C or PDB4 D. The invention also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The oligonucleotides are useful for reducing or inhibiting expression of a gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDB4 A, PDB4 B, PDB4 C, or PDB4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CCR1; CCR3; Botaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension;
                                                                                                                                                                                                                                                                                 Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADO45363 standard;
                                                                                                                                                                        The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target
                                                                                                                                                                                                                                                                                                                                                                                  Nyce JW, Sandrasagra A, Ta
Shahabuddin S, Lu H, Cong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NYCE/)
(SAND/)
(TANG/)
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23-APR-2002;
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                                                                                                                                                                                                                                        Claim 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                 CONG/)
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MILLER S.
SHAHABUDDIN
LU H.
CONG H.
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                                                                                                                                                                                                                                        SEQ ID NO 729; 174pp; English.
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2002WO-US013143.
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ng H;
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AAZ18411/c
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Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 respiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A receptor(s), and/or asthma and/or lung allergies associated with inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary disease (COPD), allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the
                                        The invention identifies a genetic locus ASTH1, associated with asthma, mapped to human chromosome 11p. ASTH11 and ASTH11 are genes present within the locus, located close to each other on human chromosome 11p, and have similar patterns of expression, and common sequence motifs. The ASTH1 genes and fragments, encoded protein, genomic regulatory regions and anti-ASTH1 antibodies are useful in the identification of individuals predisposed to development of asthma, and for the modulation of gene activity in vivo for prophylactic and therapeutic purposes. The ASTH1 protein is useful as an immunogen to raise specific antibodies, in drug screening for compositions that mimic or modulate ASTH1 activity or expression, including altered forms of ASTH1 protein, and as a therapeutic. Sequences AAZ18366-Z18509 represent polymorphisms in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ18411 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                       Mammalian asthma related genes, useful for diagnosis of a predisposition to development of asthma.
                                                                                                                                                                                                                                                                                                                                       WPI; 1999-479058/40
                                                                                                                                                                                                                                                                                                                                                                        Miller
                                                                                                                                                                                                                                                                                                                                                                                      Brooks-Wilson AR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 therapeutic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ18411;
                                                                                                                                                                                                                                                        Disclosure; Page 62; 195pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ASTH1; asthma; human; chromosome 11p;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polymorphic fragment in region 5' to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention.
                                                                                                                                                                                                                                                                                                                                                                                                                      (-SYXA)
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                              and ASTH1J genes
                                                                                                                                                                                                                                                                                                                                                                                                                      AXYS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                                                                                                                                       North M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 4 A; 3 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunogen; polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     98WO-US001260
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                                                                                                                                                                                                                                                                                                                                                                                    Buckler A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                    Cardon L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ASTH1J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ASTH11; ASTHIJ; genetic locus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                      Carey AH,
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                                                                                     family of transcription factors, which have been implicated in the activation of a variety of genes including the TCRa gene and cytokine genes known to be important in the actiology of asthma. Both ASTHII and ASTHII mRNAs are alternatively spliced. Alternative splicing of transcripts has no effect on the open reading frame of ASTHIU, as the exons involved are all 5' to the start codon in exon b. In contrast, alternative splicing of ASTHII transcripts results in 3 different ASTHII isoforms. The invention also encompasses mouse asthlj protein. The ASTHI nucleic acids are useful as diagnostics to identify a hereditary predisposition to asthma, as probes for identifying ASTHI related genes, for identifying expression of the gene in a biological specimen, and for generating genetically modified non-human animals or site specific gene modifications in cell lines. The encoded ASTHI proteins are useful as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to the ASTH1 locus on the short arm of human chromosome (1D). This locus comprises the ASTH1I and ASTH1J genes, which are associated with a genetic predisposition to asthma and bronchial hyperreactivity. The ASTH1I and ASTH1J genes are oriented in opposite directions with the ASTH1 locus, and have similar patterns of expression and common sequence motifs. They are both expressed in trachea, lung and several other tissues. ASTH1I and ASTH1J are novel members of the ets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  bronchial hyperreactivity; ets family; transcription factor; splice variant; genetic predisposition; polymorphism; antibody; drug screening; prophylaxis; therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ASTH1 protein, for e.g. s or function of ASTH1 prot predisposition to asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New nucleic acids other than naturally occurring chromosomes ASTH1 protein, for e.g. screening compositions that modulate or function of ASTH1 proteins or as diagnostics for genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-JAN-1997;
01-JUL-1997;
                                  immunogens to raise specific antibodies; in drug screening compositions that mimic or modulate activity or expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-505109/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Brooks-Wilson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-JAN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         single nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-NOV-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (AXYS-) AXYS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          187
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Miller A, North son AR, Carey AH;
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97US-0051432P.
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Pred. No. 1.4e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cardon L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Buckler A;
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                                     of ASTH1I
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therapeutic. The ASTH1 genes or fragments thereof, encoded proteins, ASTH1 genomic regulatory regions, and anti-ASTH1 and anti-ASTH11 and anti-ASTH11 and inti-ASTH11 and inti-ASTH11 and inti-ASTH11 and inti-ASTH11 and intibodies are useful in the identification of individuals predisposed to development of asthma, and for modulation of gene activity in vivo for prophylactic and therapeutic purposes. The intact ASTH11 or ASTH11 proteins or active fragments thereof may be used to modulate or reduce bronchial hyperreactivity. Sequences AAA80260-A80261 and AAA80264-A80416 represent polymorphic sites within the ASTH11 or ASTH11 genes

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Sequence 21 BP; 10 A; 7 C; 2 G; 1 T; 0 U; 1 Other;

DB 1;

0

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RESULT 951
AAH4003
ID AAH400
XX AAH400
XX AAH400
XX SINGLE
CX SNP SE
XX Single
KW SNPE;
KW SNPE;
KW Lesch-
KW polycy
KW acute
KW inflar
XX Homo:
YX WO200:
XX WO200:
XX New 9
PT 13-OC
XX New 9
PT acid
XX New 9
P
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Best Local S
Matches 18
                           primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TGGAGTTTCTCCATGTTGGT 206
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      83pp; English
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      diabetes insipidus, Lesch-Nyhan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90.0%;
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          syndrome, muscular
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The present sequence is that of single nucleotide polymorphism (SNP) probe MPO/A. The probe has the rhodamine dye dR6G at its 5' end and nitrothiazole blue (NTB) at its 3' end. It was used in a multiplex endpoint SNP analysis that demonstrated the use of novel non-fluorescent asymmetric cyanide dye compounds of the invention (NTB in the present case) as quenching reporter dyes. A 7-colour homogeneous detection of

Example 4; Col 66; 62pp; English.

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ABA91975
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 5 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Single nucleot hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Single nucleotide polymorphism probe MPO/A.
                                                                                                                                                                                                                 New non-fluorescent asymmetric cyanide dye compounds, useful quenching reporter dyes in nucleic acid hybridization assays fluorescence energy transfer as means of detection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-JAN-1998;
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                                                                                                                                                                                                                                                                                                                                                                    Lee LG,
                                                                                                                                                                                                                                                                                                                                                                                                                   (PEKE )
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                                                                                                                                                                                                                                                                                                                                                                    Graham RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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RESULT 953
AAZ87585
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Best Local S
Matches 18
The invention provides nucleic acid markers of prostate, breast and bladder cancer. The markers are indicators of malignant transformation of prostate, breast and bladder tissues and are diagnostic of the potential for metastatic spread of malignant prostate tumours. The nucleic acid can also be used as targets for therapeutic intervention in prostate cancer, benign prostatic hyperplasia (BPH), bladder cancer or breast cancer. The markers may be used to design specific probes and primers, for the rapid analysis of prostate, bladder or breast biopsy samples. The probes and primers may also be used for in situ hybridization or in situ PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid marker; biomarker; tumour; benign prostatic hyperplasia; BPH; breas diagnosis; PCR primer; ss.
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                                                                                                                                                                                                                                                                                 Example 5; Page 112; 191pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   An G,
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                                                                                                                                                                                                                                                                                                                                                                     RNA biomarkers for diagnosis, prognosis and management of prostate,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   O'hara SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Δ.
                                                                                                                                                                                                                                                                                                                                           and bladder cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rccaccreccreaecere 386
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 3 A; 10 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-00097199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99WO-US013151
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ralph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.8%;
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                                                                                                                                                                                                                                                                                    English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Veltri RW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disease marker UC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 18;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BPH; breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Band
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          #28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of the fluorogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunodetection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                bladder cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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RESULT 954
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XX Biomar
XX Prosta
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Homo s
XX US6218
XX An G,
XX C US621
CC US621
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CC Libra:
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CC C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-JUL-1995;
11-JAN-1996;
31-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           detection and diagnosis. They may also be used to identify and isolate full length gene sequences form various DNA libraries. Antibodies against the polypeptide products of the markers can be used to treat prostate cancer, bladder cancer or breast cancer. The encoded proteins may be used to detect antibodies. The proteins and antibodies can be used in immunodetection methods for detecting or quantifying the cancers, and for clinical diagnosis of these cancers. The antibodies may also be used for
                                                                                                                                                             that specifically hybridise to prostate cancer, benign prostatic hyperplasia (BPH), bladder cancer or breast cancer markers. Proteins hyperplasia (the mucleic acid markers can be used to produce antibodies ferenceded by the mucleic acid markers can be used to produce antibodies for the detection of prostate, breast or bladder cancer. The nucleic acids can be used as targets for therapeutic intervention in these diseases, the identification and isolation of full-length gene sequences, including the identification and isolation of full-length gene sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diagnosing, prognosing, and bladder cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                        detecting
                                                                                 regulatory elements for gene expression, from genomic human DNA
libraries, as hybridisation probes for screening genomic human DNA
libraries. The kits comprising the nucleic acid sequences are useful for
                                                                                                                                                                                                                                                                                                                       The sequence represents nucleic acid biomarker, UC band 28, 3' used in detection of prostate, breast and bladder cancer. Biomanucleic acid sequences can be used as hybridisation probes and
                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; Col 73; 78pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acids as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-289849/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US6218529-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Biomarker UC band 28, 3' primer #2 used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAS04002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAS04002 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  radioimaging to quantify and localize the encoded proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Prostate; breast;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UROC-) UROCOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prostate; breast; bladder; cancer; biomarker; probe; diagnostic;
benign prostatic hyperplasia; BPH; therapeutic; human; primer; a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                [2-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             O'hara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ഗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CCTCCCAAAGTGCTGGGA 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CCTCCCAAAGTGCTGGGA 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22 BP; 5
                                                        bladder,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     96US-0013611P.
96US-00692787.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95US-0001655P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-00097199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; 9 C; 5 G; 3 T; 0
                                                           breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ralph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       biomarkers and targets useful for ag, and in developing treatments for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22
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Pred. No.
                                                        prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Veltri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                           cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                           cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                      UC band 28, 3' prancer. Biomarker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human; primer; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   detecting
                                                        biological sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         prostate,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                  primer,
                                                                                                                                                                                                                                                                                                                                primers
                                                                                                                                                                seases, in including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            breast
                                                                                                                                                                                                                                                    for
                                                                                                                                                                                                                                                                                                                                                                                     #2
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Matches

18;

Conservative

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Mismatches

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Gaps

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Query Match Best Local Similarity

1.8%;

Score 18; Pred. No.

DB 1; Le 1.5e+03;

Length 22;

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RESULT 955
RAB51456
ID AAD3144
AC AAD314
AC CELL.
PA (STRA)
AC AAD314
AC AAD
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밁
                                             S
                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                Matches
                                                                                                                                                                                                                                                                           The present invention relates to methods for distinguishing between individuals homozygous for and therefore afflicted with Van Buchem's disease, individuals heterozygous for and therefore carriers of Van Buchem's disease and individuals who are not afflicted with Van Buchem's disease comprise identifying a large genomic deletion in chromosome 17 at 17q21. The method is useful for identifying individuals who are afflicted with or carriers of diseases associated with one or more genomic deletion, particularly Van Buchem's disease, which is a rare autosomal recessive disorder that results in a bone dysplasia referred to a craniotubular hypertosis. The present sequence is a PCR primer used to amplify 92Kb gene fragment in human chromosome 17 at 17q21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; Van Buchem's disease; genomic deletion; craniotubular hypertosis; autosomal recessive disorder; chromosome 17; chromosome 17q21; bone dysplasia; 92Kb gene fragment; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Methods for identifying subjects who are afflicted with or carriers of diseases associated with genomic deletion(s), e.g. Van Buchem's disease, by determining the presence of a deletion in the 92 kb region of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-JUL-2000;
06-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JUL-2001; 2001WO-US023968.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             autosomal recessione dysplasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAD31456 standard;
                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; Page 26; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        chromosome 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-227089/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Brunkow ME,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      383 CCTCCCAAAGTGCTGGGA 400
                                                   945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chromosome 17 92Kb gene fragment amplifying PCR primer,
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4
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STRAEHLING
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                                                         CAGGCTGGAGTGCAATGG
                                                                                                                                                                                                                             22
                                                                                                                                                                                                                             BP; 4
                                                                                                             1.8%;
larity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Proll S,
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2001US-0303386P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene
                                                                                                                                                                                                                             ω
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                                                                                                                                                                                                                             10 G;
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   18
                                                         962
                                                                                                                0
                                                                                                                Score 18; DB 1;
Pred. No. 1.5e+03
0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ₽,
                                                                                                                                                                                                                             5 T; 0 U; 0 Other;
                                                                                                                                                                       Length 22;
                                                                                                                      Indels
                                                                                                                   0
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                                                                                                                   Gaps
                                                                                                                   0
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RESULT 956

ABX95026/c
ID ABX950
XX
AC ABX950
XX
DT 06-JUN

ABX95026 standard; DNA;

19

0

06-JUN-2003 ABX95026;

(first entry)

RESULT 957

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ABX93650/c
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                                                                                                                                                                                                                                   The invention relates to DNA Amplification by polymerase chain reaction (PCR), comprising an artificial chromosome or a large DNA fragment of 50-5000 base pairs in length as a template and an Alu-specific primer, in 50-1000 base pairs in length as a template and an Alu-specific primer, in 50-1000 base pairs in length as a template and an Alu-specific primer, in 50-1000 base pairs in length as a template and an Alu-specific primer, in 50-1000 base pairs in 10-1000 base pairs in 10-1000 base pairs in 10-100 base pairs in 10-1000 base pairs in 10-
                                                                                    Matches
                                                                                                         Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; ss; PCR; primer; Alu repeat sequence; artificial chromosome genome chip; genetic disease; pre-labour diagnosis; tumour typing; radioactive ray damage; environmental damage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eliminating genomic repeat sequences, useful for preparing genome chips from artificial chromosomes for use in diagnosis of e.g. genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human Alu-specific 3' PCR primer Alu-N2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABX93650;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABX93650 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-268207/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Guan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-JUL-2001; 2001WO-CN001208.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-JUL-2001; 2001WO-CN001208.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003014384-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-JUN-2003
                                                                                                                                                                       Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 5; Page 8; 18pp; Chinese.
                                                                                                                                                                                                                 sequence-specific PCR primer for performing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYHK-)
                                         645
  19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             UNIV HONG KONG.
                                                                                                         Similarity
CAGGCTGRAGTGCARTGGY 1
                                           CAGGCTGGAGTGCAGTGGC 663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR; primer; Alu repeat sequence; artificial chromosome;
                                                                                                                                                                         BP;
                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first
                                                                                                                                                                         3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                         1.8%;
                                                                                                                                                                            7 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19
                                                                                                                                                                              ω
                                                                                                                                                                            <u>ن</u>
                                                                                       <u>د</u>
                                                                                                                                Score 17.8;
                                                                                                           Pred.
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                                                                                         Mismatches
                                                                                                                                                                              Τ,
                                                                                                                No. 1.4e+03
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                                                                                                                                                                                3 Other;
                                                                                                                                  DB 1;
                                                                                                                                                                                                                            method
                                                                                                                                  Length
                                                                                         Indels
                                                                                                                                                                                                                            of the invention
                                                                                         0,
                                                                                         Gaps
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AAQ757

ID AAQ757

XX AAQ757

AC AAQ757

XC AAQ757

XX O4-AUG

XX O4-AUG

XX Analys

KW Analys

KW Aggreg

XX Synthe

XX Synthe
                                                                                                                                                                                                                                                                                                           S
                                                                                                                                                                                                                                                                                                                                                                                                          밁
                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a method of amplification by polymerase chain Cr reaction (PCR) is by using an artificial chromosome or a large DNA Cr fragment of 50-5000 base pairs in length as template and an Alu-specific primer. Also included is a method for preparing a fluorescence-labelling probe comprising obtaining a polymucleotide product by performing the PCR amplification and fluorescence-labelling the polymucleotide product to give the probe. The method is useful for eliminating a repeat sequence in a genome, which is applicable in preparing genome chips from artificial chromosome for use in diagnosis of genetic diseases, pre-labour diagnosis by screening genetic diseases in pregnant women, tumour typing, diagnosis and prognosis tests and studying damages of radioactive rays and other environmental factors on humans. With this method, FISH (fluorescence inste hybridisation) probes can be produced with elimination of the Alu repeat sequence and enhanced accuracy by effectively reducing non-compared the human Alu specific PCR primer Alu-N2
                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                genome
FISH; 1
   JP06303997-A
                                Synthetic.
                                                                                                          Reverse transcription
                                                                                                                                          04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 5; Page 8; 18pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel method for eliminating repeat sequence in genome, applicable in preparing FISH (fluorescence in-site hybridization) probes from artificial chromosome for use in diagnosis of e.g. genetic diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-248303/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Guan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human Alu specific
                                                              aggregate;
                                                                              Analysis;
                                                                                                                                                                         AAQ75729
                                                                                                                                                                                                       AAQ75729 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-JUL-2001; 2001WO-CN001209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-JUL-2001; 2001WO-CN001209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYHK-)
                                                                                                                                                                                                                                                                                                              645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; ss; PCR; primer; Alu; repeat seque chip; pre-labour diagnosis; tumour fluorescence in-situ hybridisation.
                                                                                                                                                                                                                                                                                 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             UNIV HONG KONG.
                                                                                                                                                                                                                                                                                                  CAGGCTGGAGTGCAGTGGC 663
                                                                                                                                                                                                                                                                                 CAGGCTGRAGTGCARTGGY 1
                                                                             gene
                                                              restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                       BP; 3 A; 7 C; 3 G; 3 T; 0 U; 3 Other;
                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                        (first
                                                                             expression; reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer Alu-N2
                                                                                                                                                                                                       DNA;
                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                          1.8%;
                                                                                                          primer used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chinese.
                                                                                                                                                                                                       21
                                                                                                                                                                                                       ₽P
                                                                                                                                                                                                                                                                                                                                           <u>.</u>
                                                                                                                                                                                                                                                                                                                                                            Score 17.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               t sequence; fluorescence-labelling; tumour typing; radioactive ray dam
                                                                             transcription;
                                                                                                           cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                          DB 1;
.4e+03;
                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                        Length 19;
                                                                             primer; cDNA
                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                           <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ray damage;
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RESULT 959
AAQ75720
ID AAQ757
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Best Local S
Matches 19
A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of abelled reverse transcription primers (GENESEQ files AAO75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 2 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 8; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Analysis of cDNA and gene expression - by amplification of by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1994
                                                                                                                                                         Disclosure; Page 8; 11pp;
                                                                                                                                                                                                             Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                   WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                      16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-NOV-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JP06303997-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Analysis; gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Reverse transcription primer used in cDNA analysis technique
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ75720;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ75720 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                     16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19;
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                                                                                                                                                                                                                                                                                                                                  NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TTTTTATTTTTATTTTTAATT 615
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                      93JP-00112515.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.8%;
                                                                                                                                                         Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 17.8; DB 1;
Pred. No. 1.4e+03;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                    amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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                                                                                ဇ္ဇ ဇ္
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RESULT 960
AAQ97449/c
ID AAQ974
XX AAQ974
XX AAQ974
XX AAQ974
XX PCR am
XW PCR am
XW Large
XW EP6694
XX Synthe
XX Synthe
XX EP6694
XX Cheng
XX HOFF
XX Cheng
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                និនិនិនិនិ
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                                         á
                                                                                                      Matches
                                                                                                                            Query Match
Best Local :
                                                                                                                                                                                                                                                   A set of primers (Q97448-Q97455) was designed to enable the PCR amplification of the human beta-globin gene cluster. A fixed downstream primer was paired with a series of upstream primers that amplify a region extending upstream across the delta-globin gene and into the second intron of the A-gamma globulin gene. Targets of 13.5, 17.7, 19.6 and 22 kb were amplified from total human genomic DNA. A new method was used to amplify the large genomic sequences in which Thermus thermophilus DNA polymerase was used in combination with a second DNA polymerase from Thermococcus litoralis, Pyrococcus sp. or Thermatoga maritima. The present sequence (primer RHJ020) is an upstream primer corresp. to nucleotides 52152-52172 of the human beta-globin gene cluster (Genbank Acc.No. J00179) and has a Tm of 63 deg.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EP669401-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR amplification; thermostable DNA polymerase; combination; large fragment; genomic mapping; sequence analysis; beta-globin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-FEB-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-FEB-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ97449 standard;
                                                                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amplification of long nucleic Thermus thermophilus and pref.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 beta-globin gene cluster
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                                                                                                                               Similarity
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                                                    CCTGAGTAGCTGGGACTACAG
                                                                                                                                                                                                         21 BP; 4 A; 7 C; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 13; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 3 A; 0 C; 1 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.8%;
ilarity 90.5%;
Conservative
                                                                                                      Conservative
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                                                                                                                            1.8%;
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                                                                                                                            Score 17.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21
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                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              acid sequences - using a combination of . Thermococcus litoralis DNA polymerase.
                                                                                                                                 .4e+03
                                                                                                                                                        DB 1;
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RESULT 961
AAX830,4/c
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Best Local S
Matches 19
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29-DEC-1995;
30-JAN-1996;
30-JAN-1996;
12-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Primers AAX83008-X83064 were used to RT-PCR amplify exons from the 3' ends of the human WRN gene (AAX83003) which encodes a protein to Werner's syndrome. The products can be used for the detection treatment of Werner's syndrome (WS), an autosomal recessive disor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-DEC-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; WRN; Werner's syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer G to isolate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1997-363671/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oshima
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    recessive disorder; phenotype;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Isolated nucleic acid molecule encoding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   complex phenotype, as well as related diseases
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                                                                                                                                                                                                                                                                                                                                                                                        GCAGTGGTGTCATCATAGCTC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treatment of Werner's
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95US-00580539.
96US-0010835P.
96US-00594242.
96US-00632175.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6 C; 5 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 17.8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    detection; diagnosis; autosomal;
primer; RT-PCR; amplification; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RN gene product - useful and related diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
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Short tandem repeat loci; D3S1539; D4S2368; D5S818; D7S820; D9S930, D10S1239; D13S317; D14S118; D14S548; D14S562; D16S490; D16S539; D1 D17S1298; D17S1299; D19S233; D2S6481; D22S683; HUMCSF1PO; HUMTPOX; HUMTHO1; HUMFESEPS; HUMF13A01; HUMBFXIII; HUMLIPOL; HUMFWBA31; multiplex amplification reaction; MAR; allele; detection; genetic n linkage map; identification; disease gene; PCR primer; amplify; ss

detection; genetic marker;

Primer used when one of the loci in the MAR set is

D14S548

D9S930; 539; D16S753;

AAV06188;

AAV06188

standard;

21

20-MAY-1998

(first

entry)

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0222222222222200
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AAV05254
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC determining the alleles present in short tandem repeat loci from one or CC determining the alleles present in short tandem repeat loci from one or CC more DNA samples. The DNA samples to be analysed has a set of at least CC consisting of D301539, D482368, D58818, D78820, D98930, D1081239, CC CONSISTING OF D301539, D482368, D58818, D78820, D98930, D1081239, CC D138317, D148518, D148548, D148563, HUMCSF1DD, HUMTPOX, HUMTHO1, CC HUMFESFPS, HUMF13A01, HUMBFXIII, HUMLIPOL and HUWWFPA31. Alternatively, CC the DNA sample to be analysed has a set of three short tandem repeat loci which can be amplified together, where the set of loci is selected from CC which can be amplified together, where the set of loci is selected from CC the DNA sample to be sets: (1) D38139, D198253, D13817; (2) D1081239, CC D98930, D269481; (3) D1081239, D482368, D208481; (3) D1081239, D482368, D208481; (3) D1081239, D482368, D208481; D1081239, D78230, D138317; The CC loci are co-amplified in a multiplex amplification reaction (MAR), where the product of the reaction is a mixture of amplified alleles from each cof the co-amplified loci in the set. The amplified alleles in the mixture can analysed in the set within the DNA sample. The methods are used for the CC are evaluated to determine the alleles present at each of the loci can be set within the DNA sample. The methods are used for the CC detection of short tandem repeats as genetic markers for the development cof linkage maps, the identification and characterisation of disease cC genes, and the simplification and precision of DNA typing
                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Simultaneous amplification of short tandem repeats - used to provide genetic markers for linkage maps, for identifying and characterising diseases genes and for DNA typing.
                 Synthetic
                                                                   BRCAl gene; identification; mutation; multiplex amplification process; ovarian cancer; breast cancer; large scale diagnostic screening;
                                                                                                                           Sense primer used to amplify part of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21
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Homo sapiens.
                                                                                                                                                                  18-MAY-1998
                                                                                                                                                                                                                                          AAV05254 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (PROM-) PROMEGA CORP.
                                                    PCR primer;
                                                                                                                                                                                                                                                                                                                                                                        928
                                                                                                                                                                                                                                                                                                                                     21 AGRÉTEACTETETECCEAGG 1
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                                                                                                                                                                                                                                                                                                                                                                        AATCTCACTCTGTTACCCAGG 948
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 6 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                    amplify; ds
                                                                                                                                                                (first
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                                                                                                                                                                entry)
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                                                                                                                                                                                                                                          ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                            Score 17.8; DB 1;
Pred. No. 1.4e+03;
                                                                                                                           exon
                                                                                                                           21 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                           BRCA1
                                                                                                                           gene
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Homo sapiens

Lifton RP,

Simon DB

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primers AAV05254-55 are used to amplify a region of exon 21 of the BRCA1 gene. A fragment of 167 bp is produced. The primers are used in a method for identifying mutations in the BRCA1 gene using a multiplex amplification process. Mutations in BRCA1 are associated with ovarian and breast cancer. A sample is tested for mutations in the BRCA1 gene by amplifying at least one (partial) exon of the gene, and comparing the sizes and amounts of amplification products with corresponding products of the wild-type gene. Amy differences indicate a mutation. If no mutations are detected, the sequence of at least one exon may be determined. This method is inexpensive enough to be used for large scale
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detecting mutation(s) in the BRCA1 gene by exon amplification comparing amplification products with those from wild type ger optionally followed by sequencing.
                                                                                                                                                                                                       Thiazide-sensitive Na-Cl cotransporter; TSC; hTSC gene; human; ion transport; Gitelman's syndrome; Bartter's syndrome; hypokalaemic alkalosis; hypocalciuria; hypomagnesemia; diagnos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 4 A; 6 C; 6 G; 5 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diagnostic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shipman R,
                                                 31-DEC-1996;
                                                                                                     09-JUL-1998
                                                                                                                                                                                            therapy; SSCP;
                                                                                                                                                                                                                                                            Human TSC gene exon 16 reverse primer hTSCex16.
                                                                                                                                                                                                                                                                                        21-DEC-1998 (first entry)
                                                                                                                                                                                                                                                                                                                  AAV40598;
                                                                                                                                                                                                                                                                                                                                            AAV40598 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 15; Page 15; 65pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-MAY-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-NOV-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9743441-A1
                                                                            19-DEC-1997;
                                                                                                                               WO9829431-A1
                                                                                                                                                         Omo
                     (UYYA ) UNIV YALE
                                                                                                                                                       sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                     483 CAGTGGTGTGATCACAGCTCA 503
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19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                              primer;
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                                                  96US-00778052
                                                                            97WO-US023553
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Thiazide sensitive cotransporter and ATP sensitive potassium channel genes - useful for developing products for the diagnosis and treatment of ion transport disorders, e.g. Gitelman's Syndrome or Bartter's Syndrome.

Example 1; Page 51; 105pp; English. Primers hTSCex16 forward and reverse

(see AAV40597 and AAV40598

WPI; 1998-388029/33

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RESULT 965
AAZZ60
XX AAZZ60
XX AAZZ60
XX ODT 30-NOV
XX POlymc
KW Polymc
KW cell v
Cell v
W graft
XX dysplal
XX WO9841
XX WO9841
XX CO-MAP
PF 19-MAP
XX CO-MAP
XX WO1;
Ident:
PT Greves
PT dysplal
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       respectively) are designed to amplify exon 16 of the human hTSC gene (see AAV40561) that codes for thiazide-sensitive Na-Cl cotransporter TSC (see AAW29622). Both primers are located within introns of hTSC. 27 Sets of specific primers (see AAV40565-V40618) were used for SSCP analysis of hTSC. Amplified products were analysed for molecular variants by electrophoresis, and identified variants were sequenced. Complete linkage of Gitelman's syndrome with TSC was demonstrated. Identification of the molecular basis of Gitelman's syndrome allows for the genetic diagnosis of this disorder. The invention provides products and methods useful for diagnosis and treatment of Gitelman's syndrome and other ion transport
                                                                                                                                                                                                                                                                                                                                                              Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ26013 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                   Housman
                                                                                                                                                                                                            20-MAR-1997;
                                                                                                                                                                                                                                           19-MAR-1998;
                                                                                                                                                                                                                                                                        24-SEP-1998
                                                                                                                                                                                                                                                                                                    WO9841648-A2
                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human polymorphic region 202.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-NOV-1999
                                                           prevention
dysplastic
                                                        Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                (VARI-) VARIAGENICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    863 TGCTGGGATTACAGGCGTGAG 883
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                             Fig
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                                                                                                                                                  Ledley FD,
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                                                                                                                                                                                                            97US-0041057P
                                                                                                                                                                                                                                          98WO-US005419
                               7; 605pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
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90.5%;
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                                                                                                                                                   Stanton
                               English.
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Pred. No. 1.
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This invention

describes a novel method for identifying an inhibitor

Best Local Query Match

Similarity

1.8%;

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RESULT 966
AAX30235/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC on a gene vital for treatment of cancer, where the inhibitor is active CC on a gene vital for cell growth or viability, and where the gene is gubject to loss of heterozygosity (LOH) in a cancer. The inhibitor is CC used for preventing the development of cancer in a patient having a CC precancerous condition, by administering to the patient a first allele CC precancerous condition, by administering to the patient a first allele creater in cells of the precancerous condition, where the normal somatic CC cells of the patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene CC present in a population and targets only one allelic forms of the gene CC used in the diagnosis, prevention and treatment of LOH disorders, e.g. CC cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic CC lesions, benign tumours, endometriosis, polycystic kidney disease, and CC graft versus host disease. The method can also be used to remove compared to the man polymorphic sites described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer;
polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 4 A; 6 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                     region in a genetic locus using bracketing locus compatible or specificalibrating markers. The method can be used to determine DNA fragment lengths of a polymorphic region (PR) of a genetic locus, especially containing short tandem repeats. AAX30221 to AAX30248 represent PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR amplification primer b-F13A01 fwd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX30235;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX30235
                                                                                                                                                 A method
                                                                                                                                                                                Example
                                                                                                                                                                                                               Detection of length of polymorphic region in genomic loci
                                                                                                                                                                                                                                             WPI; 1999-254401/21.
                                                                                                                                                                                                                                                                          Dau PC,
                                                                                                                                                                                                                                                                                                                                      18-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                       17-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9914371-A1
                                                                                                                                                                                                                                                                                                       (OLIG-)
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                                                                                                                                                                                                                                                                                                           OLIGOTRAIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                          used
                                                                                                                                                                                3; Page 15; 63pp; English.
                                                                                                                                                                                                                                                                            Liu
                                                                                                                                                 has been developed of detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGGCTGGTCGCGAACTCCTGA 21
                                             21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amplification; bracketing; locus; electrophoresis; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 region; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                             ₽P;
                                                                          in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                            exemplification of the present
                                           6 C; 4 G; 4 T;
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Pred. No. 1.4e+03
Score 17.8; DB 1;
Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                             0 U;
                                              0 Other;
                                                                                                                                      the length of a polymorphic ng locus compatible or speci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
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              Length
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                                                                             invention
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                                                                                                                                         specific
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AAA47201-307 are oligonucleotide primers used to amplify human genomic CC DNA short tandem repeat (STR) loci. The claimed method comprises CC simultaneous determination of the alleles present in a set of loci from cone or more DNA samples. In particular, at least thirteen loci of genomic CC DNA are amplified in a single multiplex reaction. At least one of the CC loci is preferably a STR locus with a repeat unit of five to seven bases CC or base pairs in length. Preferred loci are thirteen human STR loci CC chosen by the United States Federal Bureau of Investigation as core loci for use in the Combined DNA Index System (CODIS) database. These loci are D3S1538, HUWTHO1, D2IS11, D18S51, HUWWRA31, D8S1179, HUWTPOX, HUWFIBRA, CC amplified include pentanucleotide STR loci G475, C221 and S159 (see CC AAA47308-10). Loci with intermediate length repeats can be amplified with intermediate length repeats can be amplified with CC comprises; (a) obtaining at least one DNA sample; (b) selecting a set of loci of the DNA sample comprising at least 13 short tandem repeats loci complified amplified alleles from each of the co-amplified loci in the set in a multiplex amplified alleles from each of the co-amplified loci in the set in a continue the action reaction, the product of the reaction comprising a present at each loci. The method can be used to determine the alleles of contildren, confirm the lineage of animals and agricultural crops. It is samples found at a crime scene
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AAA47233/c
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 Sequence 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New method for analyzing e.g. human tissue DNA samples comprises amplification of at least 13 short tandem repeat loci, useful in determining the parentage of a child.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Schumm JW, Sprecher CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               profile; D14S648; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer; short tandem repeat; STR; multiplex amplification reaction; Combined DNA Index System; CODIS; paternity test; breeding; forensi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Primer 1 for human genomic DNA polymorphic STR locus D14S648.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA47233;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA47233 standard;
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BP; 6 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       e.g.
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Query Match Best Local S Matches 19

Local Similarity

Conservative

0;

Mismatches

Indels

0

Gaps

0

AAF17435 standard; DNA; 21 BP

0

1.8%;

Score 17.8; DB 1 Pred. No. 1.4e+03

DB 1; Length 21;

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AAF17435/
ID AAF1
XX
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AAI70307/c
ID AAI703
                                                   RESULT 969
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                                                                                                                                                                                                                                                                            The invention provides a DNA polymerase composition for the PCR CC amplification of long (over 10 kb) nucleic acid sequences. The amplification of long (over 10 kb) nucleic acid sequences. The CC composition includes the DNA polymerase of Thermus thermophilus and a second, thermostable, DNA polymerase that provides 3'-to-5' exonuclease cC activity. Use of the composition was demonstrated for the amplification regions of the human beta-globin gene cluster, as a model for genomic trargets that are likely to contain repetitive sequences and homologous sites elsewhere in the genome. The second DNA polymerase was provided by Thermotoga maritima. Primers were designed such that a fixed downstream cc primer (see AAI70312-13) could be used with a series of upstream primers (see AAI70306-11), including the present primer, RH1020, which corresponds to nucleotides 52152-52172 of the human beta-globin gene cc cluster. Targets of 7.5-22 kb were amplified. The target region extended CC upstream across the delta-globin gene and into the second intron of the A camma globin gene. Use of primer RH1020, which lies within an Alu repeat cc sequence, resulted in multiple secondary products
                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New DNA polymerase composition consisting of a combination of a first polymerase and a second DNA polymerase, useful for amplifying nucleic acids, particularly long nucleic acid sequences by PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cheng
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-FEB-1994;
16-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-SEP-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA polymerase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-JAN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAI70307;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAI70307 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 13; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-640282/74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human beta-globin gene PCR primer RH102019.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HOFF ) HOFFMANN LA ROCHE & CO
                                                                                                                                                                                            Match 1.8%;
Local Similarity 90.5%;
                                                                                                                                          725 CCTGAGTAGCTGGGACTACAG 745
                                                                                                         21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AATCTCACTCTGTTACCCAGG 948
                                                                                                                                                                                                                                                21 BP; 4 A; 7 C; 5 G; 5 T; 0 U;
                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001EP-00113936
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94US-00203198.
95EP-00102141.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human; beta-globin; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21
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                                                                                                                                                                              0
                                                                                                                                                                                            Score 17.8; DB 1; Length 21; Pred. No. 1.4e+03;
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                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'n
                                                                                                                                                                                                                                                  0 Other;
                                                                                                                                                                               Indels
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DNA

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                                                                                                                                                                                                                                                       RESULT 970
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Best Local S
                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to DNA for a promoter and an L1 cassette sequence having a core retrotransposon element. The invention is useful for random insertion of a heterologous or homologous DNA sequence into cell genome, and for correction of a genetic defect in the cell into which the insertion is made. Genetic defects which may be corrected includes cystic fibrosis, mutations in the dystrophin gene, genetic defects associated with blood clotting and other genetic defects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Retrotransposon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              L1 cleavage site related sequence #25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF17435;
                                                                                                 Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-060015/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Moran
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15-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US6150160-A
                                                                                                                                                  SNP specific lower PCR primer SEQ ID 1202.
                                                                                                                                                                             14-AUG-2001
                                                                                                                                                                                                                               AAH38406 standard;
                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 14; 87pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 homologous DNA sequence into a cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNAc comprising a promoter P and an retrotransposon element, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYJO ) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-NOV-2000
                                                              polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; s
                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                      483
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                                                                                                                                                                                                                                                                                                                                                 19;
                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                           CAGTGGTGTGATCACAGCTCA 503
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                                                                                                                                                                                                                                                                                             CAGTGGTGATCTTAGCTCA 1
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                                                                                                                                                                                                                                                                                                                                                                                               BP; 7 A;
                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PENNSYLVANIA.
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                                                                                                                                                                             (first entry)
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96US-00749805
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genetic defect; cystic fibrosis;
                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                         1.8%;
                                                                                                                                                                                                                                                                                                                                                                                               6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                               21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kazazian HH,
                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                      Score 17.8;
                                                                                                                                                                                                                                                                                                                                                           Pred.
                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 L1 cassette sequence having a core random insertion of a heterologous genome and for correcting genetic
                                                                                                                                                                                                                                                                                                                                                           1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Boeke
                                                                                                                                                                                                                                                                                                                                                                         DB 1;
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                                                                                                                                                                                                                                                                                                                                                                      Length 21;
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ARBSULT 971
AAH37597/c
ID AAH375
XX AAH375
XX AAH375
XX II 4-AUG
DT 14-AUG
DT SNP sp
XX Single
KW SNPE;
KW SNPE;
KW Lesch
KW SUCCE
KW Inflam
XX III 1000 8
XX W OCCOOL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention CC includes kits for determining the presence or absence of a SNP, using the CC oligonucleotides of the invention. The PCR primers are used to amplify a CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or CC identity of a SNP and for genotyping nucleic acid sample by endividuals, having a pathological phenotypic trait suspected of being CC caused by one or more SNPs. Phenotypic traits include diseases e.g. CC agammaglobulinaemia, diabetes insipidus, Lesch-Myhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial cdiseases, including, rheumatoid arthritis, multiple sclerosis, include matching, rheumatoid arthritis, multiple sclerosis, and concorganism. The method is also useful in forensic investigations and contermity analysis. The present sequence represents a PCR primer specific for a human SND contration nNA securence.
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                                                                            Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rhaumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 56; 83pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   acid sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-290930/30
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                                                                                                                                                                                                           SNP specific upper PCR primer SEQ ID 393.
                                                                                                                                                                                                                                                   14-AUG-2001
                                                                                                                                                                                                                                                                                         AAH37597;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    for a human SNP containing DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1006 GATTCTCCTGTCTCAGCCTCC 1026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      l Similarity
19; Conserv
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                                                                                                                                                                                                                                                                                                                               standard;
                                                                                                                                                                                                                                                                                                                                                                                                                              GATTCTCCTGCCTCAGCTTCC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                               DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English.
                                                                                                                                                                                                                                                                                                                               21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 17.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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WO200129262-A2 Homo sapiens.

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CC She oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The coligonucleotides are useful for genotyping a nucleic acid sample by coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or coligonucleotides by association analysis the genotypic trait suspected of being constant of the present suspected of being consistency of the present suspected of being constant of the present suspected of being constant and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial consists and includes symptoms of or susceptibility to multifactorial consists and including, rheumatoid arthritis, multiple sclerosis, pathogenic mitroorganism. The method is also useful in forensic investigations and conformation, cancer, nervous system diseases and infection by pathogenic mitroorganism. The method is also useful in forensic investigations and conformation of the present sequence represents a PCR primer specific for a human SNP containing DNA sequence
AAD31450/c
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                                                                                                                                                                                                                                                                                                                                                                                           RESULT 972
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-OCT-1999;
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                                                                                                                         Human; Van Buchem's disease; genomic deletion; craniotubular hypertosis; autosomal recessive disorder; chromosome 17; chromosome 17q21; bone dysplasia; 92Kb gene fragment; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 7 A; 3 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 52; 83pp; English.
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                                                                                                                                                                                                            Human chromosome 17 92Kb gene fragment amplifying PCR primer, SpanlR.
                                                                                                                                                                                                                                                           31-MAY-2002
                                                                                                                                                                                                                                                                                                                                                 AAD31450 standard; DNA; 21 BP
                                          WO200210455-A2
                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ORCH-) ORCHID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
19; Conserv
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                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 17.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 21; .4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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07-FEB-2002

03-DEC-2001; 2001WO-US047235.

08-AUG-2002.

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RESULT 973
ABS60196/c
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to methods for distinguishing between individuals homozygous for and therefore afflicted with Van Buchem's disease, individuals heterozygous for and therefore carriers of Van Buchem's disease and individuals who are not afflicted with Van Buchems's disease comprise identifying a large genomic deletion in chromosome 17 at 17q21. The method is useful for identifying individuals who are afflicted with or carriers of diseases associated with one or more genomic deletion, particularly Van Buchem's disease, which is a rare autosomal recessive disorder that results in a bone dysplasia referred to a craniotubular hypertosis. The present sequence is a PCR primer used to amplify 92Kb gene fragment in human chromosome 17 at 17q21
                                                                                                                                               Aminopeptidase P; XPNEP2; bradykinin receptor B1; ds; BDKRB1; tachykinin receptor B1; TACRI; C1 esterase inhibitor; C1NH; kallikrein 1; KLK1; bradykinin receptor B2; BDKRB2; gene therapy; ANGI; bradykinin receptor B2; BCKRB2; gene therapy; angiotensin converting enzyme 2; ACE2; protease inhibitor 4; P14; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma; cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; aneurysm; embolism; thrombosis; coronary artery disease; angioedaema; arteriosclerosis; atherosclerosis; hypersensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Methods for identifying subjects who are afflicted with or carriers of diseases associated with genomic deletion(s), e.g. Van Buchem's disease by determining the presence of a deletion in the 92 kb region of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABS60196 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Brunkow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-JUL-2000;
06-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                              Human polymorphism associated DNA sequence #90.
                                                                                                                                                                                                                                                                                                                                                                                                  05-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                    ABS60196;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chromosome 17 at 17q21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JUL-2001; 2001WO-US023968
                                                                  WO200261131-A2
                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (STRA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7; Page 26; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21 GACCTTGTGATCCGCCCGCCT 1
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STRAEHLING HAMPTON K.
                                                                                                                                    obstructive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GACCTTGTGATCTGCCTGCCT 849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21 BP; 5 A; 5 C; 9 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                    pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Paeper B;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 17.8;
Pred. No. 1
                                                                                                                                    disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                      enterocolitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 21;
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CC tachnykinin receptor Bi (FACKI), CI esterase inhibitor (CIMI), knallikering CC (ACE2) or protease inhibitor 4 (PI4), comprising at least one CC (ACE2) or protease inhibitor 4 (PI4), comprising at least one CC (ACE2) or protease inhibitor 4 (PI4), comprising at least one CC (ACE2) or position as provided in the detailed summary of single CC polymorphic position as provided in the detailed summary of single CC equence; (2) analysing (MI) at least one nucleic acid sample comprising cobtaining the sample from one or more individuals and determining the control of haplotypes using the genes comprising grouping at least two nucleic acids expendence at one or more polymorphic positions in a gene CC encoding a protein selected from the group above; (3) constructing (M2) (M2) an individual at risk of developing a disorder CC using the polymorphic data; (5) a library of nucleic acids, each of which comprises one or more polymorphic positions within a gene encoding a comprise one or more polymorphic positions within a gene encoding a nucleic decide present in at least one polymorphic position, and (6) genotyping (M4) an individual acceptable of the control of the co
                  Best
Matches
                                      Query Match
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23-JAN-2001; 2001US-0263678P
02-MAR-2001; 2001US-0273037P
                                                                                                                                               polynucleotides are also useful for chromosome identification. Antibodies against the proteins may be utilised for immunophenotyping of cell lines and biological samples. The present sequence is included in the sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        encoding aminopeptidase P (XPNE tachykinin receptor B1 (TACR1),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 713; 977pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated nucleic acid with at least one polymorphic position, useful for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-619265/66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tsuchihashi Z, Hui L, Swanson BN, Powell JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BRIM )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            autoimmune diseases.
                                                                                    Sequence
               Local
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l Similarity 90.
19; Conservative
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TSUCHIHASHI Z.
                                                                                      21
                                                                                      BP;
                                                                                                                                 is not
                                                                                      4 A;
                                                                                                                               ples. The present sequence is incl
referred to anywhere else in the
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                                                                                      5 C;
                  1.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (XPNEP2), bradykinin receptor B1 (BI
ACR1), C1 esterase inhibitor (C1NH),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        isolated
0
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                    Score 17.8;
Pred. No. 1
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  Mismatches
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                                                                                      0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleic acid from a human gene
bradykinin receptor B1 (BDKRB1),
                      .4e+03
                                           DB 1;
                                           Length 21
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  Indels
                                                                                                                                 specification
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GGCCTCCCAAAGTGCTGGGAT 871

GGCCTCCCAAAGTACTGAGAT 1

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Хü

ABQ74069

standard; DNA; 21

ΒP

ABS98161 ID ABS RESULT 975

ABS98161 standard; DNA;

21

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0

RESULT 974 ABQ74069

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                                                                                                                                           The present invention describes a method for producing homozygous stem CC (HS) cells having a target genotype and/or immunotype from non-fertilised CC post-meiosis I diploid germ cells by mitotically activating the germ CC cells to develop multiple blastocyst-like masses, each of which contains CC an inner cell mass (ICM) that is homozygous for the target genotype CC and/or immunotype. The methods of the present invention are useful for CC the production of HS cells utilised for diagnosis, therapeutic and CC cosmetic transplantation, cell replacement and/or gene therapy, and the CC disease, Parkinson's disease (cystic fibrosis, muscular CC disease, Parkinson's disease and multiple sclerosis), traumatic injuries (post-trauma repair and reconstruction, limb replacement, spinal cord CC injuries and burns), cancer, disorders of the epithelium (Dlindness, CC diseases and anaemia. ABG/74028 to ABG/7415 represent PCR primers and CC sequence specific oligonucleotide (SSO) probes which are used in the CC exemplification of the present invention
                                         Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human leukocyte antigen; immunotype; genotype; microsatellite; probe; germ cell; nootropic; neuroprotective; antiparkinsonian; vulnerary; cytostatic; antiarteriosclerotic; antiinflammatory; immunosuppressive; antianaemic; antiidiabetic; tranquilliser; respiratory; cardiant; trauma; muscular; ophthalmological; gene therapy; genetic disease; cancer; cystic fibrosis; muscular dystrophy; cardiac condition; burn; myopathy; neurodegenerative disease; Alzheimer's disease; Parkinson's disease; muscular dystrophy; cardiac condition; burn; myopathy; neurodegenerative disease; Alzheimer's disease; Parkinson's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     producing homozygous stem cells having a target genotype and/or immunotype from non-fertilized post-meiosis I diploid germ cells suitable for diagnostic, therapeutic and cosmetic transplant and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      multiple sclerosis; post-trauma repair; reconstruction; blindness; limb replacement; spinal cord injury; atherosclerosis; Crohn's disdiabetes; autoimmune disease; anaemia; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homozygous stem cell; major histocompatibility complex; MHC; HLA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Microsatellite typing and sequencing D6S105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABQ74069;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Fig 7; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      treatment of various disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-575456/61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yan WL
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                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (STEM-)
         867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STEMRON INC
                                                            Similarity
GGGATTACAGGCGTGAGCCAC
                                                                                                                 21
                                                                                                                 BP;
                                           Conservative
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                                                                             Score 17.8;
                                                            Pred.
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                                             Mismatches
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                                                                No.
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U;
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                                                                                                                     0 Other;
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                                                                                 1;
                                                                             Length
                                                 Indels
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This invention relates to the sequence of an isolated nucleic acid conclude comprising at least one base variation from that of a known condition of the comprising at least one base variation from that of a known control of the con
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human multidrug resistance
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-698522/75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Guida M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 22; Page 144; 714pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disorder-related traits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-NOV-2000; 2000US-00724389
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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RESULT 976
ABS981
XX
ABS981
XX
ABS981
XX
ABS981
XX
ABS981
XX
DT 23-DEC
XX
DE Human
XX
Human
XX
Human
XX
Cytcoh
KW shoph
KW WaDPH
KW UDP-1
KW UDP-1
KW UDP-1
KW UDP-1
KW UDP-1
KW WADPH
KW Single
KW Centra
KW Single
KW Centra
KW Single
KW Gulda
XX
TI Solat
PT Geg. C
PT Gislat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               KW Cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1;
KW Cytochrome P450 A2; CYP4501A2; Cytochrome P450 02E; CYP45002E1; LTF;
KW dytochrome P450 A2; CYP4501A2; Cytochrome P450 02E; CYP45002E1; LTF;
KW adrenergic receptor beta1; ADBR1; aryl hydrocarbon; ARN; MRP3; NR112;
KW aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;
KW cyclooxgenase 2; COX2; diazepam binding inhibitor; DB1; haematological;
KW epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;
KW epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;
KW glutathione-5-transferase 12; GST12; histamine-N-methyl transferase; NNMT;
KW NADPH quinone oxidoreductase 2; NQO2; sulfotransferase; NNMT;
KW NADPH quinone oxidoreductase 214; UDP-glucuronosyl transferase 2B7;
KW UTP-glucuronosyl transferase 2D4; UDP-glucuronosyl transferase 2B7;
KW WDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;
KW multidrug resistance 1; lactotransferrin; orphan nuclear receptor;
KW multidrug resistance associated protein 3; cancer; prostate;
KW multidrug resistance associated protein 3; cancer; prostate;
KW acterial nervous system; pulmonary; immunological; SNP;
KW central nervous system; pulmonary; immunological; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABS98168;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200257410-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human multidrug resistance gene polymorphic sequence #70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polymorphic DNA sequence of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 1 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polymorphism.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 17.8;
Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 21;
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Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for disorder-related traits.

WPI; 2002-698522/75

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Hall J;

28-NOV-2000; 2000US-00724389

DNA SCI LAB

INC

28-NOV-2001; 2001WO-US044838

Example 22; 714pp; English

CC Cytochrome P450 (2E1 (CIP45004E1), andremerylo receptor nuclear translocator (CC (ARNT), cathepsin (ALHE), aryl hydrocarbon receptor nuclear translocator (CC (ARNT), cathepsin S (CTSS), cyclooxgenase 2 (COX2), diazepam binding CC inhibitor (DBI), epoxide hydroxylase 2 (EPHX2), 5-lipoxygenase activating CC protein (FLAP), glutathione-5-transferase 12 (CGT12), histamine-N-methyl CC transferase (NNMT), (Rallikrein 2) KLK2, nicotinamide -N-methyl CC transferase (NNMT), NADPH quinone oxidoreductase 2 (MOO2), CC (MCR1), UDP-glucuronosyl transferase 2B4 (UGT2B4), Jactotransferrin (LTF), multidrug resistance associated protein 3 (CC (MDR1), lactotransferrin (LTF), multidrug resistance associated protein 3 (CC (MDR1), lactotransferrin (LTF), multidrug resistance associated protein 3 (CC (MRR3), orphan nuclear receptor (WR12), or acetylcholine muscarinic CC receptor 1, 2, 3, 4, or 5 (CHMR1, CHMR2, CHMR3, CHMR3, CHMR3) sequence. CC (MRR3), orphan nuclear receptor (WR12), or acetylcholine muscarinic CC traits within the genome and eventually CC (MRR3), orphan muclear responsible for a variety of disorder-related CC armityling the genes responsible for a variety of disorder-related CC expression, mutation or underexpression, which may be used in diagnosing CC (ARM), multidrug resistance of their e.g., overexpression, constitutive comprising the contained in CYP4501A1, CYP94501A2, CYP4502E1, CC (MR1), MDR1 and/or MDR3 are useful for screening individuals for altered drug CC (MDR1) and MDR1 and/or MDR3 may also be used to screen individuals for altered drug CC (MDR1) and MDR1 and MDR1 and MDR1 and MDR1 and MDR1 and MDR1 and CHMR2 are cc (MDR1) and MDR2 are destine function, in CHMR3, CHMR4 or CHMR5 for altered central and contained in prous system function. The prostate in the for altered central and contained in activation in the prostate, in LTF for al molecule comprising at least one base variation from that of a known human cytochrome P450 A1 (CYP4501A1), cytochrome P450 A2 (CYP4501A2), cytochrome P450 02E1 (CYP4500E1), adrenergic receptor beta1 (ADBR1), This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known control of the comprising at least one base variation from the control of the co polymorphic DNA sequence of the invention

Sequence 21 BP; 4 A; 10 C; 4 G; 3 T; 0 U; 0 Other;

Matches 880 TGAGCCACCACGCCCGGCTTA 900 l Similarity
19; Conserv Conservative 1.8%; 0; Score 17.8; Pred. No. 1 Mismatches .4e+03; DB 1; 2 Length 21; Indels 0 0

RESULT 977 ABS98106 standard; DNA; 21

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μ,

TGAGCCACCACGCCCGTCCTA

21

(first entry)

Human multidrug resistance gene polymorphic sequence #8

ABS98106/c
ID ABS981
XX
AC ABS981
XX
AC ABS981
XX
DE Human
XX
Human;
XW Human;
XW eytoch
KW aryl |
KW aryl |
KW eycloc
KW aryl |
KW eycloc
KW apoxi,
KW Gluta;
KW GLUTa;
KW HANT;
KW HANT;
KW HANDPH
KW UDF-09
KW UGT2B Human; ds; cytochrome P450 Al; CYP4501Al; UGT2B4; MDR1; cytochrome P450 A2; CYP4501A2; cytochrome P450 02B; CYP45002B1; ITF; adrenergic receptor betal; ADBR1; aryl hydrocarbon; AHR; MRP3; NR112; aryl hydrocarbon receptor nuclear translocator; ARW7; cathepsin S; CTSS; cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological; epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP; glutathione-S-transferase 12; GST12; histamine-N-methyl transferase; HNWT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNWT; NADPH quinone oxidoreductase 2; GOX2; sulfotransferase thermolabile; STM; UDP-glucuronosyl transferase 2B7; UDP-glucuronosyl transferase 2B7; UTP-glucuronosyl transferase 2B lactotransferrin;

> multidrug resistance associated protein 3; cancer; prostate; acetylcholine muscarinic receptor; CHMR1; CHMR2; CHMR3; CHMR4; CHMR5; altered drug metabolism; cardiovascular function; colorectal tumour; central nervous system; pulmonary; immunological; SNP; Guida M, 28-NOV-2000; 2000US-00724389 WO200257410-A2 single nucleotide 28-NOV-2001; 2001WO-US044838. (DNAS-) DNA SCI LAB polymorphism

Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for disorder-related traits.

22; Page 143; 714pp; English

C expression, mutation or underexpression, which may be used in diagnosing cond/or treating the disorders. The nucleic acid molecules comprising the polymorphic sequences contained in CYP4501A1, CYP94501A2, CYP4502E1, CYP4501A1, CYP94501A2, CYP4501A2, C transferase (HNMT), (kallikrein 2) KLK2, nicotinamide -N-methyl transferase (HNMT), waDPH quinone oxidoreductase 2 (MQO2), sulfotransferase thermolabile (STM), UDP-glucuronosyl transferase 2B4 (UGT2B4), UDP-glucuronosyl transferase 2B7 (UGT2B7), UDP-glucuronosyl transferase 2B7 (UGT2B7), UDP-glucuronosyl transferase 2B7 (UGT2B7), UDP-glucuronosyl transferase 2B7 (UGT2B7), multidrug resistance 1 (MRDR), lactotransferrin (LTF), multidrug resistance associated protein 3 (MRDR), orphan nuclear receptor (NR112), or acetylcholine muscarinic receptor 1, 2, 3, 4, or 5 (CHMR1, CHMR2, CHMR3, CHMR4 or CHMR5) sequence. The polymorphisms in the human genes cited in the invention are useful as genetic linkage markers for locating and characterising the genes that are responsible for specific traits within the genome and eventually identifying the genes responsible for a variety of disorder-related traits as a result of their e.g., overexpression, constitutive aryl hydrocarbon (AHR), aryl hydrocarbon receptor nuclear translocator (ARNT), cathepsin S (CTSS), cyclooxgenase 2 (COX2), diazepam binding inhibitor (DBI), epoxide hydroxylase 2 (EPHX2), 5-lipoxygenase activating protein (FLAP), glutathione-S-transferase 12 (GST12), histamine-N-methyl transferase (HNMT), (kallikrein 2) KLK2, nicotinamide -N-methyl This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known human cytochrome P450 A1 (CYP4501A1), cytochrome P450 A2 (CYP4501A2), cytochrome P450 02E1 (CYP45002E1), adrenergic receptor betal (ADBR1) ous system function. sequence of the inve

Sequence 21 BP; տ P ហ ü 10 G 1 T; 0 U; 0 Other;

Matches l Similarity 19; Conserv Conservative 90.5%; 0 Score 17.8; DB 1; Length 21; Pred. No. 1.4e+03; Indels 0; Gaps 0

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KW cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP45002E1; LTF;
KW adrenergic receptor beta1; ADBR1; aryl hydrocarbon; AHR; MRP3; NR112;
KW aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;
KW cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological;
KW epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;
KW epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;
KW glutathione-5-transferase 12; GST12; histamine-N-methyl transferase;
KW HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT;
KW NADPH quinone oxidoreductase 21; NOO2; sulfotransferase thermolabile; STM;
KW WDP-glucuronosyl transferase 2B4; UDP-glucuronosyl transferase 2B7;
KW WDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;
KW WILTIGUT resistance 1; lactotransferrin; orphan nuclear receptor; uPA;
KW multidrug resistance 1; lactotransferrin; orphan nuclear receptor;
KW multidrug resistance associated protein 3; cancer; prostate;
MW actylcholine muscarinic receptor; CHMR1; CHMR2; CHMR3; CHMR4; CHMR5;
KW altered drug metabolism; cardiovascular function; colorectal tumour;
KW central nervous system; pulmonary; immunological; SNP;
This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known thuman cytochrome P450 A1 (CYP4501A1), cytochrome P450 02E1 (CYP4502E1), adrenergic receptor beta1 (ADBR1), cytochrome P450 02E1 (CYP4502E1), adrenergic receptor beta1 (ADBR1), cyclooxy adrenergic receptor beta1 (ADBR1), cyclooxy adrenergic receptor nuclear translocator (ARNT), cathepsin S (CTSS), cyclooxy asse 2 (COX2), diazepam binding inhibitor (DEI), epoxide hydroxylase 2 (EPHX2), 5-lipoxy genase activating protein (FLAP), glutathione-S-transferase 12 (CST12), histamine-N-methyl transferase (NNMT), (kallikrein 2) KLK2, nicotinamide -N-methyl transferase (NNMT), NADPH quinone oxidoreductase 2 (NQO2), sulfotransferase thermolabile (STM), UDP-glucuronosyl transferase 2B4 (UGT2B4), UDP-glucuronosyl transferase 2B7 (UGT2B7), UDP-glucuronosyl transferase (UGT2B15), urokinase receptor (UPA), multidrug resistance 1 (MRP3), orphan nuclear receptor (NRT2), or acetylcholine muscarinic receptor 1, 2, 3, 4, or 5 (CHMR1, CHMR2, CHMR3, CHMR4 or CHMR5) sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-698522/75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Guida M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human CYP4501A2 promoter polymorphism #2
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 102; 714pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disorder-related traits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (DNAS-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleotide polymorphism.
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RESULT 979
ABS97655
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                                                                                                                                                                                                                                                                                                                                                                                                                                         KW Cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1;
KW Cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP4502E1; LTF;
KW adrenergic receptor beta1; ADBR1; aryl hydrocarbon; AHR; MRP3; NR112;
KW aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;
KW cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological;
KW epoxide hydroxylase 2; EPHX2; S-lipoxygenase activating protein; FLAP;
KW glutathione-S-transferase 12; GST12; histamine-N-methyl transferase;
KW HNNT; kallikrain 2; KLK2; nicotinamide-N-methyl transferase; NNMT;
KW NADPH quinone oxidoreductase 2; NOO2; sulfotransferase thermolabile; STM;
KW WDP-glucuronosyl transferase 2B4; UDD-glucuronosyl transferase 2B7;
KW WT2B7; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;
KW multidrug resistance 1; lactotransferrin; orphan nuclear receptor; uPA;
KW multidrug resistance associated protein 3; cancer; prostate;
KW multidrug resistance associated protein 3; cancer; prostate;
KW acetylcholine muscarinic receptor; uPA;
KW multidrug muscarinic receptor; upa;
KW acetylcholine muscarinic receptor; upa;
KW acetylcholine muscarinic receptor; upa;
KW central nervous system; pulmonary; immunological; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression, mutation or underexpression, which may be used in diagnosing carding or treating the disorders. The nucleic acid molecules comprising the polymorphic sequences contained in CYP4501A1, CYP4502B1, CYP4502B1, CYP4502B1, CYP4502B1, CYP4502B1, CYP4502B1, CYP4502B1, CYP4501A1, CYP4501A2, CY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABS97655
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                                                                                                                28-NOV-2000; 2000US-00724389
                                                                                                                                                                            28-NOV-2001;
                                                                                                                                                                                                                                                                                                   WO200257410-A2
                                                                                                                                                                                                                                                                                                                                                                                                                   single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human glutathione-S-transferase 12 (GST12) polymorphic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polymorphic DNA sequence of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TGATCCGCCCGTCTCGGCCTC 21
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                                                                                                                                                                               2001WO-US044838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                      polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 17.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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Guida M,

(DNAS-) DNA SCI LAB INC

Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for

disorder-related traits.

2002-698522/75

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RESULT 980
ABA99519/c
                                                                                                                                                                     밁
                                                                                                               S
                                                                                                                            Matches
                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                      This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known human cytochrome P450 A1 (CYP4501A1), cytochrome P450 A2 (CYP4501A2),
                                                                                                                                                                                                                                                                                                                                                                                                                Example 12; Page 122; 714pp; English
                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                 cytochrome P450 02E1 (CYP45002E1), adrenergic receptor betal (ADBR1),
    Tumour-associated antigen; human; B345; cytostatic; cell communication;
cell interaction; signal transduction; metastasis; cancer; colon;
immunotherapy; carcinoma; lung; diagnosis; PCR; primer; ss.
                              Human tumour-associated antigen B345 PCR primer SEQ ID NO
                                                                    ABA99519
                                                                                                                374
                                                                                                                            19;
                                                                                                    ۳
                                                                                                                                  Similarity
                                                                                                                CTGCCTCAGCCTCCCAAAGTG 394
                                                                                                    CTGCCTCAGCCTCACAAAGCG
                                                                                                                                                     21
                                                                    standard;
                                                                                                                                                     BP;
                                                                                                                                                                  DNA sequence of the invention
                                           (first entry)
                                                                                                                                                     տ
                                                                                                                                                     Þ
                                                                    DNA;
                                                                                                                                 1.8%;
                                                                                                                                                     9 C; 4 G; 3 T; 0 U; 0 Other;
                                                                     21
                                                                                                                            0
                                                                                                                                  Score 17.8;
Pred. No. 1.
                                                                                                    21
                                                                                                                            Mismatches
                                                                                                                                  1.4e+03
                                                                                                                                          DB 1;
                                                                                                                                        Length 21;
                                                                                                                             Indels
                                16
                                                                                                                             0
                                                                                                                             Gaps
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0

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RESULT 981
ADH47846/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ligands, especially in the metastatic potential of cancers, particularly of the colon. B345 or its immunogenic fragments, also the DNA that encodes it, are useful for immunotherapy of cancer, particularly carcinoma of lung or colon. Antibodies raised against B345 are useful for treatment and diagnosis of cancers that are associated with B345 expression, including their use for targeted delivery of cytotoxic or radioactive agents. Probes derived from B345 can be used to detect tumour specific mutations in the B345 sequence, and can be used to screen for B345 specific modulators. This sequence represents a PCR primer used in the amplification of the human B345 tumour-associated antigen described
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes a novel tumour-associated antigen, designated B345 which has cytostatic activity. B345 is involved in communication, interaction and/or signal transduction with extracellular components and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200204508-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New tumor-associated antigen B345, useful for diagnosis and immunotherapy of tumors, also related nucleic acid and antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUL-2000;
19-APR-2001;
                                                                                     Antidiabetic; anorectic; cardiant; hypotensive; antiarteriosclerotic; anorectic; virucide; antibacterial; fungicide; protozoacide; nootropic neuroprotective; antiparkinsonian; anticonvulsant; osteopathic; antiarthritic; antiinflammintory; dermatological; antiasthmatic; antiinflammintory; human; metabolic disorder; diabetes; obes; viral infection; bacterial infection; fungal infection; helminthic infection; protozoal infection; anorexia; cancer; helminthic infection; protozoal infection; anorexia; cancer; disease; neurodegenerative disorder; hlzheimer's disease; parkinson's disease; pellepsy; immune disorder; harmopoietic disorder; harmopoietic disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-171704/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-JUL-2001; 2001WO-EP007705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-JAN-2002
                                                                                                                                                                                                                                                                              25-MAR-2004
                                                                                                                                                                                                                                                                                                                                          ADH47846 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BOEH ) BOEHRINGER INGELHEIM INT
                                                                                                                                                                                                                                                                                                           ADH47846;
                                                                        inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                    991 CTCCCGGGCTCAAGCGATTCT 1011
                                                                                                                                                                                                                                                                                                                                                                                                                       21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                       CTCCTGGGCTCAAGCAATTCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000DE-01033080
2001DE-01019294
                                                                                                                                                                                                                                               SEQ ID
                                                                                                                                                                                                                                                                              (first entry)
                                                                            skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Scherl-Mostageer M,
                                                                             disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ç
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 17.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ħ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sommergruber W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ų;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ĎΒ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>,,,</u>
                                                                                            haematopoietic disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Abseher
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                     nootropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                            disease;
                                                                                                                                                         obesity;
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RESULT 982
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                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 19
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18-JAN-2001; 2001US-026587P.
31-JAN-2001; 2001US-0265530P.
14-FEB-2001; 2001US-0268595P.
28-FEB-2001; 2001US-0276409P.
16-MAR-2001; 2001US-0276777P.
17-MAY-2001; 2001US-027677P.
17-MAY-2001; 2001US-035306P.
18-CCT-2001; 2001US-0345202P.
Human; NOV; adrenoleukodystrophy; congenital adrenal hyperplasia; haemophilia; hypercoagulation; autoimmune disease; allergy; immunodeficiency; transplantation; Von Hippel-Lindau syndrome;
                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to novel proteins (I) referred to as NOVX, where X is any number from 1 to 18, and their coding sequences (II). The proteins and their coding sequences are useful in the manufacture of a medicament for treating a syndrome associated with a human disease, preferably a NOVX-associated disorder such as metabolic disorders, diabetes, obesity, infectious diseases (viral, bacterial, fungal, helminthic, and protozoal), anorexia, cancer, cardiovascular diseases (hypertension, atherosclerosis), neurodegenerative disorders, Alzheimer's disease, Parkinson's disease, epilepsy, immune disorders (osteoarthritis), haematopoletic disorders, inflammatory skin disorders, asthma, and various dyslipidaemias. The present sequence is a probe for a NOVX sequence. This sequence has a TET modification at the 5' end and a TAMRA modification at the 3' end.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vernet CAM, Li L, Shenoy, Macdougall J, Malyankar U, Tchernev V, Zerhusen BD, Baumgartner J, Herrmann J, Taupier RJ, Gerlach V, Gr
                                                                                                                                                                                                                                                                                                                                                            Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated NOVX polypeptides and polynucleotides, useful for preventing, diagnosing or treating NOVX-associated disorders e.g. osteoarthritis, obesity, atherosclerosis, cancer, Parkinson's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-SEP-2002
                                                                Novel human
                                                                                             16-MAY-2003
                                                                                                                                                        ABX97680 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Page 346; 380pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2002-698671/75.
                                                                                                                                                                                                                                                                  646
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                                                                                                                                                                                                                                    21
                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          or infections.
                                                                                                                                                                                                                                                                 AGGCTGGAGTGCAGTGGCGCA 666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Burgess CE;
                                                                                                                                                                                                                                                                                                                                                          BP; 3
                                                              protein
                                                                                                                                                                                                                                                                                             1.8%;
llarity 90.5%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Alsobrook JP, Colman SD, Spytek KA, Boldog F;
Li L, Shenoy S, Casman S, Guo X, Edinger S;
Malyankar U, Patturajan M, Shimkets RA, Pene,
Zerhusen BD, Millett I, Miller C, Lepley DM,
J, Herrmann J, Peyman JA, Gorman L, Mezes P,
Gerlach V, Grosse WM, Liu X, Ellerman K, Rotl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001US-0261376P
2001US-0262454P
                                                                                             (first
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                                                                                                                                                                                                                                                                                                                                                            ð,
                                                                                                                                                        DNA;
                                                              NOVX associated reverse PCR primer
                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                          11 C;
                                                                                                                                                        21
                                                                                                                                                                                                                                                                                                                                                          3 G;
                                                                                                                                                        ₽₽
                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                              Score 17.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                          4 T;
                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                            0
U;
                                                                                                                                                                                                                                                                                                              .4e+03;
                                                                                                                                                                                                                                                                                                                          DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                            0 Other;
                                                                                                                                                                                                                                                                                               N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Edinger'S;
ets RA, Pena
                                                              #15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P, Kekuda
Rothenberg
                                                                                                                                                                                                                                                                                             ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Smithson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 7
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The invention describes an isolated polypeptide, NOVX, comprising a CC sequence or a mature form of one of 21 51-1543 residue amino acid CC sequences (P1-P21), given in the specification. The NOVX polypeptides, CC polynucleotides and antibodies are useful in the manufacture of a medicament for treating or preventing e.g. adrenoleukodystrophy. CC congenital adrenal hyperplasia, haemophilia, hypercoagulation, autoimmune CC disease, allergies, immunodeficiencies, transplantation, Von Hippel-CC Lindau syndrome, Alzheimer's disease, stroke, tuberous sclerosis, CC hypercalcaemia, Parkinson's disease, Huntington's disease, Cancer, CC fertility, diabetes, adult respiratory distress syndrome, viral, CC bacterial and parasitic infections. The nucleic acid sequences may be used in chromosome mapping, identifying individual from minute biological comples (tissue typing), and in forensic identification of a biological comple. This sequence represents a primer used to isolate DNA encoding a covel human protein (NOV)
                                                                                Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-MAY-2001;
09-MAY-2001;
11-MAY-2001;
11-MAY-2001;
14-MAY-2001;
15-MAY-2001;
21-MAY-2001;
23-MAY-2001;
25-MAY-2001;
29-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patturajan M,
Kekuda R, Go;
Mezes PS, Pej
                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New NOVX polypeptides and polynucleotides useful for treating preventing e.g. congenital adrenal hyperplasia, hemophilia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-MAY-2001;
07-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Alzheimer's disease; stroke; tuberous sclerosis; hypercalcaemia; Parkinson's disease; Huntington's disease; cancer; fertility; di adult respiratory distress syndrome; infection; tissue typing; forensic identification; gene; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-103511/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Spytek KA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200290500-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hypercoagulation, autoimmune disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CURA-) CURAGEN CORP.
21
                                                                                    19;
                                                                                                        Similarity
                                                                                                                                                                      21
                                        ATTACAGGCGTGAGCCACCAC 890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gorman
Peyman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 273; 300pp; English
                                                                                                                                                                      B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Li L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2001US-0288395P.
2001US-0289619P.
2001US-0289619P.
2001US-0289817P.
2001US-0289818P.
2001US-0290194P.
2001US-0290753P.
2001US-029174P.
2001US-0293747P.
2001US-02941494P.
2001US-0328446P.
2001US-031326446P.
2001US-00136728.
                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2002WO-US014256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            L, Edinger SR,
Gerlach VL, Ta
                                                                                                                                                                      ທ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ĮĄ,
                                                                                                                                                                      A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                      90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                zerhusen BD, Smithson G, Zhong M;
                                                                                                                            1.8%;
                                                                                  0;
                                                                                                        Score 17.8;
Pred. No. 1.
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                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          allergies,
                                                                                                        .4e+03
                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Guo X,
ena CEA,
                                                                                                                          Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunodeficiencies,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Macdougall
                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Anderson
Padigaru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fertility; diabetes;
                                                                                    0
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                                                                                  Gaps
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RESULT 983
ACF64055/c
RESULT 984
ACC6403
ID ACC640
XX
ACC ACC640
AC ACC640
XX
13-OCT
XX
DE IFNAR1
CX
XX
Human;
KW signal
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                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                The present invention describes an isolated polynucleotide (PN) comprising: (a) a sequence comprising at least 15 contiquous nucleotides of a sequence comprising variant sequences (A) from Table 4 given in the specification; or (b) a sequence that is complementary to (A). Also described: (1) an array of (PN)s comprising two or more of the isolated (PN)s; (2) detecting a (PN) in an individual; (3) a computer-readable storage medium, where each record has a field identifying a base occupying a (PN) site and a location of the polymorphic site; and (4) a signal carrying data for access by an application program having executed on a data processing system. The (PN) can be used for detecting loci associated with multiple sclerosis. ACF64025 to ACF64424 represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; detection; computer-readable storage medium; polymorphic site; signal carrying data; data processing system; multiple sclerosis; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-AUG-2001; 2001US-0310741P.
24-SEP-2001; 2001US-0324790P.
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                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New polynucleotide, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-268196/26.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-FEB-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IFNAR1 forward PCR primer #31
Human; detection; computer-readable storage medium; polymorphic site; signal carrying data; data processing system; multiple sclerosis;
                                                                                                                    ACF64053 standard; DNA;
                                        IFNAR1
                                                                  13-OCT-2003
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19; Conserv
                                        forward
                                                                                                                                                                                                             GCTGGAGTGCAGTGGCGCAAT 668
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                                                                                                                                                                                   GCTAGAGTGCAGTGGTGCAAT 1
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                                                                                                                                                                                                                                                                                           BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 10; 93pp;
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                                                                                                                                                                                                                                      Conservative
                                                               (first entry)
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                                        PCR
                                                                                                                                                                                                                                                                                           5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          White
                                      primer
                                                                                                                                                                                                                                                                                           8 C;
                                                                                                                                                                                                                                                 1.8%;
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                                                                                                                    21
                                                                                                                                                                                                                                                                                             3 G; 5 T; 0 U; 0 Other;
                                        #29
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                                                                                                                      ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for detecting loci associated with multiple
                                                                                                                                                                                                                                        ٥.
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                                                                                                                                                                                                                                                   Score 17.8;
Pred. No. 1.
                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                    1.4e+03
                                                                                                                                                                                                                                                                 DB 1;
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                                                                                                                                                                                                                                                                  Length 21;
                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Natsoulis G;
                                                                                                                                                                                                                                        0;
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RESULT 985
ADF11633/c
ID ADF116
XX
AC ADF116
XX
AC ALTER

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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              comprising: (a) a sequence comprising at least 15 contiguous nucleotides of a sequence comprising variant sequences (A) from Table 4 given in the specification; or (b) a sequence that is complementary to (A). Also described: (1) an array of (PN) so comprising two or more of the isolated (PN)s; (2) detecting a (PN) in an individual; (3) a computer-readable storage medium, where each record has a field identifying a base occupying a (PN) site and a location of the polymorphic site; and (4) a signal carrying data for access by an application program having executed on a data processing system. The (PN) can be used for detecting loci associated with multiple sclerosis. ACF64025 to ACF64424 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-AUG-2001; 2001US-0310741P.
24-SEP-2001; 2001US-0324790P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-AUG-2002; 2002WO-US025268.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-268196/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jones
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                                              03-APR-2002; 2002US-0370088P
                                                                                               03-APR-2003; 2003WO-US010649
                                                                                                                                                                                                                                                                                                                osteopathic;
                                                                                                                                                                                                                                                                                                                                                                   Alternate human SRP5/SRP6 polymorphism reverse primer.
                                                                                                                                                                                                                                                                                                                                                                                                                    12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF11633 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (DNAS-) DNA SCI INC
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                                                                                                                                                23-OCT-2003
                                                                                                                                                                                              WO2003087763-A2
                                                                                                                                                                                                                                             sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -ب
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity 90.5
19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACTCCCGACCTCAGATGATCC 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACTCCTGACCTCAGGTGATCC 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention describes an isolated polynucleotide (PN)
                                                                                                                                                                                                                                                                                           gene therapy; bone mineral density; sclerostin gene; osteopenia; bone dysplasia; bone fracture; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      White
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8 C; 4 G; 5 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           망
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rienhoff HY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 17.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    detecting
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2;
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(CELL-) CELLTECH R & D INC.

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RESULT 986
ADF11652/c
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     presence or absence of at least one sclerostin gene region nucleotide polymorphism in a biological sample from a subject where the presence of at least one polymorphism at a position that corresponds to a non-coding region of the 130320 by sclerostin gene region (SOST) indicates an increased risk of altered BMD. The composition and methods are useful in determining in a subject a risk for having, or presence of, altered bone mineral density, such as osteoporosis, osteopenia, bone dysplasia, bone fracture or other conditions characterized by decreased or increased bone density. These may also be used in identifying agents that may be used for treating the above diseases, disorders or conditions associated with altered BMD. In addition, these may be used for pharmacogenomic purposes, e.g. to stratify patient populations according to suitability of a particular therapeutic agent for use in the population. This sequence corresponds to the reverse primer for the alternative human sclerostin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Determining a risk for or presence of altered bone mineral density (e. osteoporosis) in a subject comprises determining the presence or abserof a sclerostin gene region nucleotide polymorphism in a biological
                                                                                                                                                                                                                                                                                                                                                                                                                              ADF11652 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 7 A; 5 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 21; 114pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-833790/77.
                                                                                                                                                                                                                                                                                        osteopathic; gene therapy; bone mineral density; sclerostin gene osteoporosis; osteopenia; bone dysplasia; bone fracture; primer;
                                                                                                                                                                                                                                                                                                                                  Human sclerostin gene region polymorphism 5 reverse primer.
                                                                                                                                                                                                                                                                                                                                                                    12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of altered
                              WPI; 2003-833790/77
                                                                                                                                    03-APR-2002; 2002US-0370088P
                                                                                                                                                                   03-APR-2003; 2003WO-US010649
                                                                                                                                                                                                                               WO2003087763-A2.
                                                                                                                                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention relates to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
                                                                                        CELLTECH R & D
UNIV ROTTERDAM
                                                            ME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          bone mineral density
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a subject
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                  (first
                                                          Charmley PR, Proll S,
                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.8%;
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                                                                                        INC.
ERASMUS
                                                                                                                                                                                                                                                                                                                                                                                                                              21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      a method of determining a risk for or prodensity (BMD) in a subject by determining
                                                                                                                                                                                                                                                                                                                                                                                                                              ₽₽
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 17.8;
Pred. No. 1
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                                                          Paeper BW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4e+03
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                                                            Uitterlinden
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               absence
                                                                                                                                                                                                                                                                                        region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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Determining a risk for or presence

of altered bone mineral density (e.g.

Creating an insertional mutation in the germ line of an animal, useful for generating a mutation in an offspring of an animal, comprises introducing into an animal a nucleic acid molecule comprising a germ l

a germ line

WPI; 2003-863454/80

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a method of determining a risk for or presence or faltered bone mineral density (BMD) in a subject by determining the presence or absence of at least one sclerostin gene region nucleotide polymorphism in a biological sample from a subject where the presence of at least one polymorphism at a position that corresponds to a non-coding region of the 130320 by sclerostin gene region (SOST) indicates an increased risk of altered BMD. The composition and methods are useful in determining in a subject a risk for having, or presence of, altered bone mineral density, such as osteoporosis, osteopenia, bone dysplasia, bone confirmed density. These may also be used in identifying agents that may be used for treating the above diseases, disorders or conditions associated with a letered BMD. In addition, these may be used for pharmacogenomic purposes, corresponds to the forward primer for the human sclerostin gene region corresponds to the forward primer for the human sclerostin gene region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local
                                                                                                                                                                       16-NOV-1995;
15-NOV-1996;
28-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    osteoporosis) in a subject comprises determining the presence or absence of a sclerostin gene region nucleotide polymorphism in a biological sample from a subject.
                                                                                                                                                                                                                                                                                                                                                                            gene therapy; insertional mutation; germ line specific promoter; non-LTR;
                                                                                                                                                                                                                                                                                         US2003121063-A1
                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                            retrotransposon;
cleavage site; ds
                                                                                                                                                                                                                                                                                                                                                                                                                      L1 retrotransposon endonuclease cleavage site seq id 116
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADF12370;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADF12370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 7 A; 5 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polymorphism 5
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                                                                                                                                                                                                                                 09-AUG-2002; 2002US-00216122.
                                                                                                                                                                                                                                                             26-JUN-2003
                                                                                                                              (UYPE-) UNIV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CGGATTCAAGTGATTCTCCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CGGGTTCAAGTTATTCTCCTG 715
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                                                                                                                              PENNSYLVANIA.
                                                                                                                                                           2000US-00653812
                                                                                                                                                                       95US-0006831P.
96US-00749805.
97US-00847844.
                                                                                                 Ostertag
                                                                                                                                                                                                                                                                                                                                                  d8
                                                                                                                                                                                                                                                                                                                                                           lon; transgenic animal; poly A element; non-LTR;
long terminal repeats; L1; EN domain; endonuclease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.8%;
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Pred. No. 1.
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Example

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116; 102pp; English

The invention describes a method of creating an insertional mutation in the germ line of an animal by introducing into an animal a nucleic acid molecule comprising a germ line specific promoter. The method is useful for generating a mutation in an offspring of an animal, or for isolating a nucleic acid from a genome of an offspring of an animal. The method may also be used to correct genetic defects in animals, especially humans. The nucleic acid is useful for generating mutations in a cell for assessing the frequency with which selected cells under go insertional mutagenesis for the generation of transgenic animals. This sequence represents an exemplary cleavage site of the endonuclease encoded by human L1 retrotransposon EN domain.

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CAGTGGTGTGATCTTAGCTCA 1 CAGTGGTGTGATCACAGCTCA 503

483 21

Query Match Best Local S Matches 19

l Similarity 19; Conserv

Conservative

<u>,</u>

1.8%;

Score 17.8; Pred. No. 1. Mismatches

DB 1; L.4e+03; 0 Other;

Length Indels

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF12525
NT 000501 contig on chromosome 8p21-22 or a polymorphic marker which is in linkage disequilibrium with the chromosome. The PCM1 marker is preferably D8S261, D8S2615 or D8S2616 and lies within the PCM1 gene. The novel method involves assessing two or more of the PCM1 markers single
                                                                                                                                                               This invention describes a novel method of determining the susceptibility to or diagnosis of schizophrenia comprising using a marker located in the chromosomal region 8p21-22. The method involves determining the presence or absence in a test sample of a pericentriolar material 1 (PCM1) marker which is selected from any of the microsatellite repeats present in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Determining the susceptibility of an individual to a neuropsychiatric disorder (e.g. schizophrenia) or diagnosing or prognosing the disorder comprises using a pericentriolar material 1 marker in the chromosomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-532919/50.
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                                                                                                                                                                                                                                                                                                                                                                                                             Claim 30; Page 67; 108pp;
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                                                                                                                                                                                                                                                                                                                                                                                                             English
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                                                               within the intronic sequence 3' to exon 4, in exon 4, or in the intronic sequence 5' of exon 5. The PCMI marker is assessed by strand conformation polymorphic marker analysis, heteroduplex analysis or restriction fragment length polymorphism (RFLP) analysis. Schizophrenia therapy comprises screening an individual for a genetic predisposition to schizophrenia, where the predisposition is correlated with the PCMI marker and if a predisposition is identified, providing therapeutic treatment for the individual. Alternatively, the method comprises administering to a patient a substance that modulates the expression from the PCMI gene or a gene located within 1000 kbase of the PCMI locus. This sequence represents a primer sequence used to detect novel microsatellite repeats identified on the PCMI D8S2616 marker found on the NT 000501
Sequence 21
BP; 6
A; 5 C; 5
G; 5 T; 0 U;
     0 Other
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Matches Query Match Best Local 385 19;  $\vdash$ Similarity TCCCAAAGTGCTGGGATTACA 405 Conservative 1.8%; 90.5%; 0 Score Pred. Mismatches e 17.8; L. No. 1.4e+03; DB 1; Length Indels 0; Gaps 0

RESULT 989 ADH59601/c Synthetic probe. non-nucleotide ADH59601 standard; DNA; 21 WO2003027328-A2 Non-nucleotide 25-MAR-2004 ADH59601; (first entry) probe; Bacterial Artificial Chromosome clone; probe of the ΒP invention BAC;

03-APR-2003 Kirtsen 24-SEP-2001; 2001US-0324499P. 24-SEP-2002; 2002WO-US030573 (DAKO-) ,VN DAKOCYTOMATION DENMARK BOSTON PROBES Hyldig-Nielsen JJ, AS Williams

Non-nucleotide probe for suppressing binding of detectable nucleic acid probes to undesired sequences, has aggregate nucleobase sequence homologous to randomly distributed repeat sequence of genomic nucleic acid SEQ ID NO 7; 103pp; English

WPI; 2003-421160/39.

Claim 10;

The present sequence represents a non-nucleotide probe. The probe is useful for suppressing the binding of one or more detectable nucleic ac probes, that are greater than 100 base pairs and that have been derived from genomic nucleic acid, to one or more undesired sequences in an ass for determining target genomic nucleic acid of a sample. The method comprises contacting the sample with the mixture of probes (preferably comprising 5-50 probes), contacting the sample with the one or more detectable nucleic acid probes, and determining the target genomic nucleic acid of the sample by determining the hybridization of the one or more detectable acid of the sample by determining the baseline of the one or more detectable acid of the sample by determining the hybridization of the one or more detectable mucleic acid of the sample by determining the hybridization of the one or more detectable mucleic acid of the sample by determining the hybridization of the one or more detectable mucleic acid probes, and determining the target genomic nucleic acid of the sample by determining the hybridization of the one or more detectable mucleic acid probes. detectable nucleic acid probes to cleic acid derived an assay e one ရှိ ရှိ

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RESULT 990
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Non-nucleotide probe for suppressing binding of detectable nucleic acid probes to undesired sequences, has aggregate nucleobase sequence homologous to randomly distributed repeat sequence of genomic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-SEP-2002; 2002WO-US030573
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            non-nucleotide probe; Bacterial Artificial Chromosome clone; BAC; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Non-nucleotide
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                                                                                                                                                                                                              Kirtsen
                                                                                                                                                                                                                                                                                                                                                                                 24-SEP-2001; 2001US-0324499P
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                                                                                                                                                                                                                                                                              (DAKO-)
                                                                                                                                                                                                                                                                                                                 (BOST-)
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                                                                                                                                       2003-421160/39
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                                                                                                                                                                                                          NV,
                                                                                                                                                                                                                                                                              DAKOCYTOMATION DENMARK AS
                                                                                                                                                                                                                                                                                                                 BOSTON PROBES INC
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                                                                                                                                                                                                          Hyldig-Nielsen JJ, Williams
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Claim 10; SEQ ID NO 19; 103pp; English

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                                                                                                                                                infectious disease; anorexia; cancer; cardiovascular disease; hypertension; atherosclerosis; neurodegenerative disorder; Alzheimer's disease; Parkinson's disease; epilepsy; immune disorder; osteoarthritis; hematopoietic disorder; inflammatory skin disorder; asthma; dyslipidemia; neurogenesis; cell differentiation; cell proliferation; hematopoiesis; wound healing; anglogenesis;
                                                                                                                                                                                                                                                                                                                                                                         intidiabetic; anorectic; cardiant; hypotensive; antiarteriosclerotic; anorectic; virucide; antibacterial; fungicide; protozoacide; nootropic; neuroprotective; antiparkinsonian; anticonvulsant; osteopathic; antiarthritic; antiinflammatory; dermatological; antiasthmatica; antilipemic; gene therapy; metabolic disorder; diabetes; obesity; antilipemic; gene therapy; metabolic disorder; diabetes; obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 4 A; 2 C; 7 G; 8 T; 0 U; 0 Other;
                                                                                                                chromosome mapping; tissue typing; pharmacogenomic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel NOVX gene sequence forward primer #39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-MAY-2004 (first entry)
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19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 17.8; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 21;
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20-NOV-2001; 2001ÜS-0331630P.
20-NOV-2001; 2001ÜS-0331641P.
21-NOV-2001; 2001US-0332152P.
28-NOV-2001; 2001US-0333402P.
29-NOV-2001; 2001US-033402P.
29-NOV-2001; 2001US-0334300P.
30-NOV-2001; 2001US-0334526P.
04-DEC-2001; 2001US-0334526P.
04-DEC-2001; 2001US-0334526P.
04-DEC-2001; 2001US-0338330P.
04-DEC-2001; 2001US-0338330P.
01-EEB-2002; 2002US-0338330P.
01-FEB-2002; 2002US-0353280P.
01-FEB-2002; 2002US-0353280P.
04-FEB-2002; 2002US-0353280P.
04-FEB-2002; 2002US-0353280P.
04-FEB-2002; 2002US-0353280P.
04-FEB-2002; 2002US-0354393P.
04-FEB-2002; 2002US-0354393P.
04-FEB-2002; 2002US-0354393P.
04-FEB-2002; 2002US-0354393P.
05-MAR-2002; 2002US-0354393P.
05-MAR-2002; 2002US-0351790P.
05-MAR-2002; 2002US-0361833P.
05-MAR-2002; 2002US-0361833P.
05-MAR-2002; 2002US-0361833P.
05-MAR-2002; 2002US-0364182P.
13-MAR-2002; 2002US-0364182P.
23-MUG-2002; 2002US-0364182P.
23-MUG-2002; 2002US-0364182P.
23-MUG-2002; 2002US-036482P.
23-MUG-2002; 2002US-036482P.
23-MUG-20
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06-NOV-2001; 2001US-0333072F
09-NOV-2001; 2001US-0348283F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-NOV-2002; 2002WO-US035464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2003040325-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-MAY-2003
                                                                                                              Agee ML, Alsobrook JP, Berghs C, Boldog FL, Burgess CE, Chant JS Chaudhuri A, Dipippo VA, Edinger SR, Eisen A, Ellerman K; Gangolli EA, Gorman L, Gerlach VL, Ji M, Kekuda R, Khramtsov NV; Li L, Malyankar UM, Macdougall JR, Mezes PS, Miller CE, Millet I Ooi CE, Ort T, Padigaru M, Patturajan M, Rastelli L, Rieger DK; Rothenberg ME, Shenoy SG, Spaderna SK, Spytek KA, Taupier RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-NOV-2001;
16-NOV-2001;
New isolated NOVX polypeptides and preventing, diagnosing or treating
                                                          WPI; 2003-441551/41
                                                                                                                                                                                                                                                   CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001US-0335610P.
2001US-0338543P.
2001US-0331630P.
2001US-0331641P.
2001US-033461P.
2001US-0333461P.
2001US-0334912P.
2001US-0334027P.
2001US-0334300P.
2001US-033426P.
2001US-0334526P.
2001US-0334526P.
                                                                                                BD,
                                                                                              Zhong
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    polynucleotides, useful for NOVX-associated disorders,
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parkinson's disease, epilepsy, immune disorders (osteoarthritis), hematopoietic disorders, inflammatory skin disorders, asthma, and various dyslipidemias. The nucleic acids and polypeptides may also be used as targets for the identification of small molecules that modulate or inhibit e.g. neurogenesis, cell differentiation, cell proliferation, hematopoiesis, wound healing and angiogenesis, in gene therapy, in generation of antibodies that bind immunospecifically to NOVX substances for use in therapeutic or diagnostic methods. The nucleic acids are further used as hybridization probes, in chromosome mapping, tissue
                                                                                                                                                                                                                                                                                                                                                                 these, or a sequence that is at least 95 % identical to, or having one of more conservative amino acid substitutions in the polypeptides. The polypeptides, nucleic acid molecules and antibodies are useful in the manufacture of a medicament for treating a syndrome associated with a human disease, preferably a NOVX-associated disorder. The nucleic acid molecules, polypeptides and antibodies are useful for treating, preventing or diagnosing diseases such metabolic disorders, diabetes, obesity, infectious diseases (viral, bacterial, fungal, helminthic, and protozoal), anorexia, cancer, cardiovascular diseases (hypertension, atherosclerosis), neurodegenerative disorders, Albeimer's disease,
                                   typing, preventive medicine, and pharmacogenomics. This sequence corresponds to a forward primer for the genes encoding one of the NC polypeptides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            osteoarthritis, obesity, atherosclerosis, cancer, Parkinson's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 567; 800pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      asthma, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    relates to novel isolated polypeptides, mature forms of equence that is at least 95 % identical to, or having one
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                                                                                     of the NOVX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ő
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밁 Ś Query Match Best Local & Matches 220 AACTCCCGACCTCAGATGATC 240 19; 1 AACTCCTGACCTCAGGTGATC Similarity Conservative 90.5%; 1.8%; <u>.</u> Score 17.8; Pred. No. 1 Mismatches .4e+03 DB 1; 2; Length 21; Indels 0; Gaps

0

Sequence 21 BP; 5 A; 7 C; 4 G; 5 T; 0 U; 0 Other;

RESULT 992 ADK01282 06-MAY-2004 (first entry) Rat DNA microarray capture oligonucleotide ADK01282; ADK01282 standard; DNA; 21 ВP #2.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus

DE10208794-A1

04-SEP-2003.

28-FEB-2002; 2002DE-01008794

28-FEB-2002; 2002DE-01008794.

JS;

DEGUSSA BIOACTIVES GMBH

Boekenkamp D, Dieck HT, норре н;

2003-714082/68

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

e.g.

Example; Page 4; 8pp; German.

This invention describes a novel method for sorting single-stranded conclusic acids by isolation and hybridisation of nucleic acid pools, then comparise that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at cleast once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture clasture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceranic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardivity of pharmaceuticals and/or nutritional compounds, e.g. food acitives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and can detect very small differences in pools of nucleic acid from cells. It can detect very small differences in the nucleic acid stom cell, and can detect very small differences in the mucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual capture probes used in the method of the invention.

Sequence 21 BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;

RESULT 993 ADK01329 TO THE PROPERTY OF THE PROPERT 밁 Ś Query Match Best Local Matches ADK01329 standard; 428 19; Similarity TTTTATTTTATTTTTTTAAG 448 Conservative DNA; 1.8%; 21 ВP <u>;</u> Score 17.8; Pred. No. 1. Mismatches 1.4e+03 DB 1; Length 21; 2 <u>,</u>

ADK01329

06-MAY-2004 (first entry)

Rat DNA microarray capture oligonucleotide #49.

ss; hybridisation; nerve; tion; capture oligonucleotide; pattern; mucosal; hair root; germ cell; food additive; food supplement. food supplement.

DE10208794-A1

28-FEB-2002; 2002DE-01008794

28-FEB-2002; 2002DE-01008794

DEGUSSA BIOACTIVES GMBH.

Boekenkamp D, Dieck HT,

WPI; 2003-714082/68

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

CC that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture care agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture cinceraction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the cativity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic carids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible can detect very small differences in pools of nucleic acids from cells. It can detect very small differences in the nucleic acids from cells, and comparison of nucleic acid pools, not individual capture probes used in the method of the invention. agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present a least once in each nucleic acid and a 3'-variable, discriminatory region This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) at

Sequence 21 BP; 2 A; 0 C; 0 G; 19 T; 0 U; 0 Other;

0

Ś Query Match Matches Local 427 TITTTATTTTATTTTTTAA 447 l Similarity
19; Conserv Conservative 1.8%; 90.5%; 0, Score 17.8; DB 1; Pred. No. Mismatches 1.4e+03 Length 21; Indels 0; Gaps 0

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ADP68377 standard; DNA;

ADP68377;

12-AUG-2004 (first entry)

DNA probe used to detect human NOV14 DNA (Ag210) SeqID 261

human; probe; ss; NOVX; Alzheimer's disease; Huntington's; inflammatory; crohn's disease; rheumatoid arthritis; immunological; endocrine; pigmentation; haematopoietic; psychotic; autoimmune; muscular; osteoporosis; angina pectoris; hypotension; anxiety; allopecia; bulimia; cancer; manic depression; virucide; antibacterial; analgesic; neuroprotective; nootropic; cerebroprotective; anticonvulsant; dermatological; osteopathic; antiarthritic; antinflammatory; cytostatic hypotensive; cardiant; hypertensive; antiulcer; antiallergic; hypotensive; hypotens antianginal; immunosuppressive; antidepressant; neurodegenerative cytostatic;

Homo sapiens.

WO200281510-A2

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RESULT 995
ADF86416/c
                                                                                                                                                                                                                                                                                                                                          This invention relates to novel nucleic acid molecules encoding NOVX CC polypeptides selected from NOV1 to NOV11 inclusive, as well as variants CC thereof. Specifically, it refers to vectors, host cells, antibodies, CC agonists, antagonists and recombinant methods for producing proteins CC including GpCRs, secretory proteins and dual specificity phosphatases. CC The present invention describes these proteins as useful for the CC diseases such as Alzheimer's and Huntington's, inflammatory conditions CC including Crohn's disease and rheumatoid arthritis, as well as CC including Crohn's disease and rheumatoid arthritis, as well as CC inmunological, endocrine, pigmentation, haematopoietic, psychotic, CC autoimmune and muscular disorders. Accordingly, it refers to various conditions including osteoporosis, angina pectoris, hypotension, anxiety, CC allopecia, bulimia, cancer and manic depression. As such, they exhibit various activities including vulnerary, virucide, antibacterial, canalogsic, neuroprotective, nootropic, cerebroprotective, anticonvulsant, dermatological, osteopathic, antiarthritic, antianflammatory, cytostatic, hypotensive, cardiant, hypertensive, antiallergic, osteopathic, antiarthritic, antiallergic, can signification of the invention.
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Best Local Similarity
Matches 19; Conser
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23-JAN-2001; 2001US-026605P.
25-JAN-2001; 2001US-0264159P.
31-JAN-2001; 2001US-026517P.
97-FEB-2001; 2001US-026598P.
27-FEB-2001; 2001US-027098P.
27-FEB-2001; 2001US-0271855P.
02-MAR-2001; 2001US-0272920P.
18-APR-2001; 2001US-0285940P.
24-APR-2001; 2001US-0286287P.
05-JUL-2001; 2001US-0303229P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anderson
Ellerman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New NOVX polypeptides useful for treating cancers, blood disorders, asthma, psoriasis, vascular disorders, hypertension, viral, bacterial parasitic infections, allergy, renal disorders and skin disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-JAN-2002;
                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CURA-) CURAGEN CORP.
             VLA4 antagonist-related PCR primer #1
                                                                                                         ADF86416
                                              26-FEB-2004
                                                                            ADF86416;
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                                                                                                                                                                                                                     646
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                                                                                                                                                                                      AGGCTGGAGGGCAGTGGTGCA 1
                                                                                                                                                                                                                     AGGCTGGAGTGCAGTGGCGCA 666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 261; 415pp; English.
                                                                                                                                                                                                                                                                                                                   BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Stone DJ,
                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Burgess CE, Casman SJ, Col-
Gerlach V, Gunther E, Keku
Patturajan M, Rothenberg M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002WO-US001467
                                              (first entry)
                                                                                                                                                                                                                                                                                                                   11 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                      90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vernet CAM,
                                                                                                                                                                                                                                                                                   . 88
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                                                                                                                                                                                                                                                                                     Score 17.8; DB 1;
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                                                                                                                                                                                                                                                       Mismatches
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Shimkets RA,
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Niiteu Y,
Tanaka I,
                                                                                                                                                                                                                                                                                                                                                                                                                                The invention comprises VLA4 antagonists that may optionally be other anticancer agents for the treatment of acute leukaemia. Tantagonists of the invention may be used to treat, prevent and acute leukaemia, the VLA4 antagonists may also be used to scree candidates. The present DNA sequence represents a PCR primer thused in an example of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VLA4, antagonist; acute leukaemia; screening; PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Treatment and/or prevention of acute leukemia with medicinal compositions containing VLA4 antagonist, also applicable in diagnosing its prognosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 5 A; 7 C; 3
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                                                                                                                                                                                                                                06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3;
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                             27-JUN-2002; 2002DK-00001005
07-OCT-2002; 2002DK-00001500
25-FEB-2003; 2003DK-00000289
29-APR-2003; 2003DK-00000639
                                                                                27-JUN-2003; 2003WO-DK000448
                                                                                                                        WO2004003229-A2
                                                                                                                                              Synthetic.
                                                                                                                                                        Homo sapiens
                                                                                                                                                                          sequence polymorphism analysis;
single nucleotide polymorphism;
                                                                                                                                                                                                           Human chromosome 19 RAI il anchor probe.
                                                                                                                                                                                                                                                                         ADK41377
                                                                                                     08-JAN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                screening drug candidates.
                                                                                                                                                                                                                                                                                                                                                 389 AAAGTGCTGGGATTACAGGCG 409
                                                                                                                                                                                                                                                                                                                              21
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MATSUNAGA
                                                                                                                                                                                                                                                                          standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matsunaga T, Takemoto N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 1; 72pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                (first entry)
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                  Pred.
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                                                                                                                                                                            SNP; probe
                                                                                                                                                                                       human; chromosome 19q;
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                                                                                                                                                                                                                                                                                                                                                                                                                Other;
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                                                                                                                                                                                                                                                                                                                                                                                          Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Akiyama
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             optionally be used with e leukaemia. The VLA4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                primer that
                                                                                                                                                                                          cancer;
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screen drug
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                                                                                                                                                                                          RAI;
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UNIV AARHUS. ARBEJDSMILJO

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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer to a disease treatment; a primer or probe for use in the method of estimating the disease risk or prognosis of an individual or for estimating a treatment response of an individual suffering from cancer to a disease treatment; an antibody directed to an epitope of a RAI gene product; and a kit for use in the method of estimating the disease risk or prognosis of an individual or for estimating a treatment response of an individual suffering from cancer to a disease treatment, comprising at least one primer or probe and optionally amplifying means for nucleic acid amplification. The novel method is useful for estimating the disease risk or prognosis of an individual or for estimating a treatment response of an individual suffering from cancer to a disease treatment. This polymorate in the sequence represents a probe used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nexo
Estimating the disease risk or prognosis of an individual by sequence
                                                                                                                                                                                                                                                 27-JUN-2002; 2002DK-00001005.
07-OCT-2002; 2002DK-00001500.
25-FEB-2003; 2003DK-00000289.
29-APR-2003; 2003DK-00000639.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence polymorphism analysis; human; chromosome 19q; cancer; RAI;
single nucleotide polymorphism; SNP; probe; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel method of estimating disease risk or prognosis of an individual by sequence polymorphism analysis, especial polymorphisms in the human chromosome 19q. The invention further relates to: estimating a treatment response of an individual suffering from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; SEQ ID NO 135; 145pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Estimating the disease risk or polymorphism analysis.
                                                     WPI; 2004-142878/14
                                                                                                                                                                                                                                                                                                                                                                                                                                               08-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human chromosome 19 DNA primer/probe SEQ ID No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADK41251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADK41251 standard;
                                                                                                                                                                                                                                                                                                                                                                                       27-JUN-2003; 2003WO-DK000448.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2004003229-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention.
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                                                                                                         Vogel U,
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                                                                                                           Rockenbauer
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Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prognosis of an individual by sequence
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                                                                                                                                                                 OCCUPA.
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                                                                                                           Bukowy ZK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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CC prognosis of an individual by sequence polymorphism analysis, especially color polymorphisms in the human chromosome 19q. The invention further relates concert to a disease treatment response of an individual suffering from concert to a disease treatment; a primer or probe for use in the method of estimating a treatment response of an individual suffering from cancer to a disease treatment; an antibody directed to an epitope of a RAI gene corproduct; and a kit for use in the method of estimating the disease risk or prognosis of an individual or for estimating a treatment response of an individual suffering from cancer to a disease treatment response of corprognosis of an individual or for estimating a treatment, comprising at least one primer or probe and optionally amplifying means for nucleic corisk or prognosis of an individual or for estimating at treatment response of an individual suffering from cancer to a disease treatment response of an individual suffering from cancer to a disease treatment response corplymucleotide sequence represents a primer/probe used for detecting the disease treatment. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 30; SEQ ID NO 9; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polymorphism analysis
21
BP; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    relates to a novel method of estimating disease risk or an individual by sequence polymorphism analysis, especia
Α,
3 C; 11 G;
3 T;
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0 Other;
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RESULT 998
ADM32266
ID ADM3222
XX ADM322
XX ADM322
XX DT 20-MAY
XX Human
XX human
XX Homo 6
OS Synthe
XX JP2004
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XX JP2007
XX JP2007
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      single nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human interleukin-18; IL-18; adult onset still disease; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-MAY-2004
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l Similarity 90.5%;
19; Conservativo
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The invention relates to a novel method for detecting a gene polymorphism in a human interleukin (II)-18 gene. The method involves detecting a 9 base insertion between -6311 position and -6310 position, a polymorphism at positions -5890, -5316, -4762, -4675, -3268, -689 and -640 of a polymorphism polymucleotide which consists of a fully defined sequence of 6640 base

Detecting gene polymorphism detecting adult onset still

in interleukin-18 gene of human, disease.

WPI; 2004-174121/17.

22-JUL-2002; 2002JP-00212550 22-JUL-2002; 2002JP-00212550

SUGIURA S.
HYUBITTO GENOMICS

JP2004049136-A Synthetic Homo sapiens

19-FEB-2004

Claim 6;

SEQ ID NO

23; 61pp; Japanese.

밁 S

Sequence

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RESULT 999
ADL25728/c
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PARCE 
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16-MAR-2001;
17-MAY-2001;
27-SEP-2001;
18-OCT-2001;
09-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-JAN-2001;
18-JAN-2001;
18-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                pairs as given in the specification, where in the 6640bp polynucleotide, the position 6575 is set to +1 from which numbering is performed. The method is useful for detecting gene polymorphism in II-18 gene of human and for detecting adult onset still disease. This polynucleotide sequence represents a primer of the human interleukin-18 gene of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vaccine; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ss; probe; Cytostatic; Neuroprotective; Immunosuppressive; Gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human NOVX gene,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-MAY-2004
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(SHEN/)
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(ALSO/)
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(BOLD/)
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14-FEB-2001;
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19; Conserv
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COLMAN S D.
SPYTEK K A.
BOLDOG F L.
VERNET C A M.
LI L.
EDINGER S R.
MACCOUGALL J R.
MALYANKAR U M.
SHIMKETS R A.
PENA C E A.
TCHERNEV V T.
ZERHISEN B D.
MILLET I.
MILLET C E.
LEPLEY D M.
SMITHSON G.
BAUMGARTNER J C.
                                                                                                                                                                                                                                                               SHENOY S G. CASMAN S J. GUO X.
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2001US-0262454P.
2001US-0262587P.
2001US-0265530P.
2001US-026559P.
2001US-0277409P.
2001US-0276777P.
2001US-0276772P.
2001US-0330336P.
2001US-0345202P.
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                                                                                                                                                                                                                                                                                                                                                                                                                      1 C1 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                 ס
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8 C; 5 G; 4 T; 0 U; 0 Other;
    a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        #29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 뫋
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 17.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ..4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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(UYRO-) (DAVI/)

UNIV DAVI

ROTTERDAM ERASMUS F B L. 2002US-0417779P.

11-OCT-2002;

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RESULT 1000
ADM94155/c
ID ADM94155 standard; DNA;
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Paddigaru M, Alsobrook JP, Colman SD, Spytek KA, Boldog FL Vernet CAM, Li L, Shenoy SG, Casman SJ, Guo X, Edinger SI, Macdougall JR, Malyankar UM, Patturajan M, Shimkets RA, Py Tchernev VT, Zerhusen BD, Millet I, Miller CE, Lepley DM; Smithaon G, Baumgartner JC, Herrmann JL, Peyman JA, Gorman Mezes PD, Kekuda R, Taupier RJ, Gerlach V, Grosse WM, Linellerman K, Rothenberg M, Stone DJ, Burgess CE;
                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to novel human NOVX nucleic acids and polypeptides. The polypeptide, nucleic acid or antibody is useful for preparing a composition for treating or preventing a NOVX-associated disorder, e.g., neurodegenerative or autoimmune disorders or cancer. The present sequence represents a probe used to isolate human NOVX genes of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New NOVX polypeptide, useful for preparing a preventing a NOVX-associated disorder, e.g., autoimmune disorders or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HERR/)
(PEYM/)
(GORM/)
(KEZE/)
(KEZE/)
(TAUP/)
(GERL/)
(GERL/)
(GROS/)
(LIUX/)
(ELLE/)
(ROTH/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-081706/08.
                                                                                                                        Synthetic
                                                                                                                                                                  nucleic acid amplification; primer; PCR; detection,
                                                                                                                                                                                       BCL-2 gene related
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 3 A; 11 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example
                                                       13-OCT-2003;
                                                                            22-APR-2004.
                                                                                                  WO2004033728-A2
                                                                                                                                             lymphoproliferative
                                                                                                                                                                                                               15-JUL-2004
                                                                                                                                                                                                                                      ADM94155;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BURG/)
                                                                                                                                                                                                                                                                                                                                         646
                                                                                                                                                                                                                                                                                                                   21 AGGCTGGAGGGCAGTGGTGCA
                                                                                                                                                                                                                                                                                                                                                                19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEZES P D.
KEKUDA R.
TAUPIER R J.
GERLACH V.
GROSSE W M.
LIU X.
ELLERMAN K.
ROTHENBERG M.
STONE D J.
BURGESS C E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HERRMANN J L.
PEYMAN J A.
GORMAN L.
                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ω
•
                                                                                                                                                                                                                                                                                                                                 AGGCTGGAGTGCAGTGGCGCA 666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 263;
                                                                                                                                                        translocation; clonal rearrangement;
                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                      2003WO-NL000690
                                                                                                                                                                                                               (first
                                                                                                                                                                                         3'MBR2
                                                                                                                                                                                                               entry)
                                                                                                                                              disorder;
                                                                                                                                                                                                                                                                                                                                                                           1.8%;
90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   282pp; English.
                                                                                                                                                                                                                                                            21
                                                                                                                                                                                         primer.
                                                                                                                                                                                                                                                            ₽P
                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                           Score 17.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                           1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     composition for treating neurodegenerative or
                                                                                                                                                                                                                                                                                                                                                                  2;
                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                          chromosome aberration;
                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Boldog FL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gorman L;
                                                                                                                                                                                                                                                                                                                                                                                         21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pena
                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CEA;
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RESULT 1001
AAT71928/c
ID AAT7192
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a set of nucleic amplification primers C capable of amplifying a VH-JH or DH-JH IGH, VK-JK or VK/intron-Kde IGK, C Vlambda-Jlambda IGL, Vbeta-Jbeta TCRB or Dbeta-CRB, Vy-JY TCRG, C Vdelta-Jdelta, Ddelta-Ddelta or Vdelta-Ddelta TCRD rearrangement C comprises a forward primer and a reverse primer. Also described: (1) a C nucleic acid amplification assay, preferably a PCR or multiplex PCR C assay, using the set of primers; (2) detecting VH-JH or DH-JH IGH, VK-JK C or VK/intron-Kde IGK, Vlambda-Jlambda IGL, Vbeta-Jbeta TCRB or Dbeta-C Jbeta TCRB, VV-JY TCRG, Vdelta-Jdelta or Vdelta-Ddelta or Vdelta-Ddelta C TCRD rearrangement; (3) detecting chromosomal translocation (11;14) (BCLg-C IGH); (4) detecting buman TBXAS1, recombination C G C TCRD rearrangement; (5) assessing clonal rearrangements and/or chromosome aberrations; and (6) a kit for the detecting at least one rearrangement C comprising the set of primers. The new set of nucleic amplification C G TCRG, Vdelta-Jdelta, Ddelta-Ddelta TCRD rearrangement C G Lambda-Jlambda IGL, Vbeta-Jbeta TCRB or VK-JK or VK/intron-C C G G IGK, Vlambda-Jlambda IGL, Vbeta-Jbeta TCRD rearrangement C G Lambda-Jlambda IGL, Vbeta-Jbeta TCRD rearrangement C G Lambda IGL, Vbeta-Jbeta TCRD rearrangement C G Lambda IGL, Vbeta-Jbeta TCRD rearrangement C G Lambda IGL, Vbeta-Jbeta TCRD Lambda IGL, Vbeta-Jbeta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Van
Garzia San
Garzia PAS,
                                                                  08-MAY-1995;
15-NOV-1995;
09-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New set of nucleic amplification primers comprising a forward primer and a reverse primer and capable of amplifying a rearrangement, useful in diagnosing lymphoproliferative disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-364878/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Primer; polymerase chain reaction; amplify; hereditary haemochromatosis; HH; mutation; HH-associated allele; base-pair polymorphism; HHP-1; HHP-19; HHP-29; microsatellite repeat allele; genetic marker; interferon treatment; hepatitis C infection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT71928 standard; DNA; 22
                                                                                                                                                                                                                                                                                    14-NOV-1996
                                                                                                                                                                                                                                                                                                                                                       WO9635803-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer detects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-AUG-1997
(MERC-) MERCATOR GENETICS INC
                                                                                                                                                                                                              08-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dongen JJM, Langerak
zia Sanz R, Parreira
ns PAS, Kneba M, Hum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1052 GCCACCACACCCCGCTAATTT 1072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fig 11A; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                  95US-00436074.
95US-00559302.
96US-00599252.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                marker 4072-2 in
                                                                                                                                                                                                              96WO-US006583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Langerak AW, Schuuring EMD, San Miquel JF;
Parreira A, Smith JL, Lavender FL, Morgan
Na M, Hummel M, Macintyre EA, Bastard C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; 3 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 17.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                臣
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Morgan GJ;
rd C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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The sequences given in AAT71901-72 represent a series of primer pairs cwhich were used to determine the presence or absence of the common cometation in an individual. The method comprises assessing genomic DNA from an individual for the polymorphism HHP-1, HHP-19 or HHP-29, and/or at least one non-optional comarker comprising the following microsatellite repeat alleles of group A common comprising the following microsatellite repeat alleles of group A compositionally of group B: Group A: 1990/205), 1884(235), 1A2(239), CC 184(271), 24E2(245), 2B8(206), 3321-1(197), 4073-1(182), 4440-1(180), CC 4440-2(139), 731-1(177), 590-1(148), 950-5(180), 950-5(180), 950-2(134), 950-3(165), 950-4(128), 950-5(180), 950-5(180), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Diagnosing and genotyping of hereditary haemochromatosis (HH) -primers to detect specific polymorphisms of the HH gene on chromogeneous control or novel microsatellite markers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Drayna DT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14; Page 14; 67pp; English.
                                                                                             this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Feder JN,
                                                                                      potential,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gnirke
                                                                                          the responsiveness of interferon treatment may
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Þ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Thomas WJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wolff
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chromosome
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밁 S Query Match Best Local S Matches 19 935 21 19; Similarity CTCTGTTACCCAGGCTGGAGT 955 CTCTATTGCCCAGGCTGGAGT Conservative 1.8%; <u>.</u> Score 17.8; Pred. No. 1. Mismatches 1.5e+03 ďB Length Indels 22; 0, Gaps 0

Sequence

22 BP; 6

₽,

7 C; 6

G; 3 T; 0 U; 0 Other;

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RESULT 1002
AAT71942/c
AAT71942;
                 AAT71942
                 standard;
                 DNA;
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18-AUG-1997

(first entry)

Primer detects marker 950-8 in HH region of chromosome

6p2.1.

08-MAY-1995; 15-NOV-1995; 09-FEB-1996; Synthetic. Primer; polymerase chain reaction; amplify; hereditary haemochromatosis; HH; mutation; HH-associated allele; base-pair polymorphism; HHP-1; HHP-19; HHP-29; microsatellite repeat allele; genetic marker; interferon treatment; hepatitis C infection; ss. 08-MAY-1996; 14-NOV-1996 WO9635803-A1. 95US-00436074. 95US-00559302. 96WO-US006583.

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RESULT 1003
AAT71925/c
ID AAT71925/c
XX AAT7192
XX AAT7192
XX AAT7192
XX Primer
XX Interfe
XX Synthet
XX Synthet
XX Primer
XX OS-MAY
XX OS-MAY
XX OS-MAY
PR 15-NOV
PR 15-NOV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The sequences given in AAT71901-72 represent a series of primer pairs cc which were used to determine the presence or absence of the common cc method comprises assessing genomic DNA from an individual. The method comprises assessing genomic DNA from an individual for the cc method comprises assessing genomic DNA from an individual for the cc presence or absence of the HH-associated allele of the base-pair cc marker comprising the following microsatellite repeat alleles of group A cc and optionally of group B: Group A: 19D9(205), 18B4(235), 1A2(239), cc 4440-2(139), 731-1(177), 5091-1(148), 3216-1(221), 4073-1(182), 4440-1(180), cc 4440-2(139), 731-1(177), 5091-1(148), 3216-1(221), 4072-2(148), 950-(151), 250-(164), 950-3(165), 63-1(128), 63-2(169), 63-3(169), 63-3(169), 65-1(206), 65-2(81), 373-8(151), cc 373-29(109), 68-1(167), 241-6(105), 241-29(113) Group B: D68364(206), cc 263(238), D68256(199), D68265(122), D68105(124) and D681001(180); cc where the number in brackets indicates the number of nucleotides between and including the flanking primers and the absence of the genotype cc indicates the likelihood of the presence of the Hutation. Knowledge of the new genetic markers allows the definition of genotypes characteristic cof heterozygous carriers and homozygotes having a HH mutation in their cc genomic DNA, The potential for HH in an individual interferes with the cffectiveness of interferon treatment for hepatitis C infection. By the first of the presence of the presence of the HT may the flanking primars of interferes with the cffectiveness of interferon treatment for hepatitis C infection. By
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Diagnosing and genotyping of hereditary haemochromatosis (HH) primers to detect specific polymorphisms of the HH gene on ch 6p2.1 or novel microsatellite markers.
 08-MAY-1995;
15-NOV-1995;
09-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                  18-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                          AAT71925 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MERC-) MERCATOR GENETICS
                                                                                                                                                                                                                                                                                                              Primer detects
                                                                           08-MAY-1996;
                                                                                                              14-NOV-1996.
                                                                                                                                                                                                                      interteron
                                                                                                                                                                                                                                           HHP-19;
                                                                                                                                                                                                                                                                           Primer;
                                                                                                                                                                                                                                   mer; polymerase chain reaction; amplify; hereditary haemochromatosis; mutation; HH-associated allele; base-pair polymorphism; HHP-1; 19; HHP-29; microsatellite repeat allele; genetic marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1996-518691/51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     931
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14; Page 15; 67pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         l Similarity
19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTCACTCTGTTACCCAGGCTG 951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTCACTCTGTCTCCCAGGCTG 1
                                                                                                                                                                                                                    treatment; hepatitis C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                  (first entry)
   95US-00436074.
95US-00559302.
96US-00599252.
                                                                                                                                                                                                                                                                                                              marker 3216-1 in HH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6 A;
                                                                           96WO-US006583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.8%;
90.5%;
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 H,
                                                                                                                                                                                                                                                                                                              region of chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Thomas WJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                               6p2.1.
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The sequences given in AAT71901-72 represent a series of primer pairs committee which were used to determine the presence or absence of the common commended comprises assessing genomic DNA from an individual. The common complete common common complete co
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6p2.1 or novel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 14; Page 14; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Drayna DT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MERC-) MERCATOR GENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and genotyping of hereditary haemochromatosis (HH) -detect specific polymorphisms of the HH gene on chronovel microsatellite markers.
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Query Match Best Local S Matches 19 Sequence 935 Similarity CTCTGTTACCCAGGCTGGAGT 955 Conservative 1.8%; 0; Score 17.8; Pred. No. 1 Mismatches H ū; 1.5e+03 DB 1; Length Indels 22; 0

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Other;

맑 RESULT 1004 AAT72000/c 21 á

AAT72000 standard; DNA; 18-AUG-1997 AAT72000 detects (first marker entry) 4072-2 22 ВP ä HH region of chromosome

6p2.1.

08-MAY-1995; 15-NOV-1995; Primer; polymerase chain reaction; amplify; hereditary haemochromatosis; HH; mutation; HH-associated allele; base-pair polymorphism; HHP-1; HHP-19; HHP-29; microsatellite repeat allele; genetic marker; 06-MAY-1996; 14-NOV-1996. interferon treatment; 95US-00436074 95US-00559302 96WO-US006352 hepatitis C infection; ss

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RESULT 1005
AAT71997/c
ID AAT7199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cc method comprises assessing genomic DNA from an individual for the CC presence or absence of the HH-associated allele of the base-pair CC polymorphism HHP-1, HHP-19 or HHP-29, and/or at least one non-optional CC marker comprising the following microsatellite repeat alleles of group A CC 184(271), 24E2(245), 2B8(206), 3321-1(197), 4073-1(182), 4440-1(180), CC 184(271), 731-1(177), 5991-1(148), 3216-1(221), 4072-2(148), 950-2(164), 950-3(165), 950-4(128), 950-5(180), 950-6(151), 950-CC 1(142), 950-2(164), 950-3(165), 950-4(128), 950-5(180), 950-6(151), 950-CC 1(142), 950-2(164), 63-1(167), 241-6(105), 65-1(201), 4072-2(148), 950-CC 1(165), 63-1(120), 63-1(619), 63-3(169), 63-1(121), 173-8(151), 273-8(151), 273-2(109), 68-1(167), 241-6(105), 241-29(113) Group B: D68464(206), CC D68306(238), D68258(199), D68265(122), D68105(124) and D681001(180); CC where the number in brackets indicates the number of nucleotides between cand including the flanking primers and the absence of the genotype CC indicates the 1kelihood of the presence of the HH mutation. Knowledge of the new genetic markers allows the definition of genotypes characteristic CC of heterozygous carriers and homozygotes having a HH mutation in their CC genomic DNA. The potential for HH in an individual interferes with the CC diagnosing this potential, the responsiveness of interferon treatment may be absence of interferon treatment may be absenced in the properties and homozygotes having a HH mutation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Determn. of the common hereditary haemochromatosis gene mutation - using primers based on novel microsatellite repeat flanking sequences or on base-pair polymorphisms HHP-1, HHP-19 or HHP-29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The sequences given in AAT71973-2044 represent a series of primer pairs which were used to determine the presence or absence of the common hereditary haemochromatosis (HH) gene mutation in an individual. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 14; Page 14; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-518690/51
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                                                                                                                                                                                                                             Primer; polymerase chain reaction; amplify; hereditary haemochromatosis; HH; mutation; HH-associated allele; base-pair polymorphism; HHP-1; HHP-29; microsatellite repeat allele; genetic marker; interferon treatment; hepatitis C infection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
08-MAY-1995;
                                                                                                                                                                                                                                                                                                                                             Primer detects marker 3216-1 in HH region of chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT71997 standard; DNA; 22 BP
                                          06-MAY-1996;
                                                                                          14-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                                        18-AUG-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                lagnosing this potential, evaluated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    935 CTCTGTTACCCAGGCTGGAGT 955
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95US-00436074
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 17.8;
Pred. No. 1
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The sequences given in AAT71973-2044 represent a series of primer pairs CC which were used to determine the presence or absence of the common CC hereditary haemochromatosis (HH) gene mutation in an individual. The CC method comprises assessing genomic DNA from an individual for the CC presence or absence of the HH-associated allele of the base-pair CC polymorphism HHP-1, HHP-19 or HHP-29, and/or at least one non-optional CC marker comprising the following microsatellite repeat alleles of group A CC 1E4(271), 24E2(245), 2B8(206), 3321-1(197), 4073-1(182), 1A2(239), CC 1E4(271), 24E2(245), 2B8(206), 3321-1(197), 4073-1(182), 4440-1(180), CC 4440-2(139), 731-1(177), 5991-1(148), 3216-1(221), 4072-2(148), 950-2(144), 950-3(165), 950-4(128), 950-5(180), 950-5(180), 950-5(180), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950-6(151), 950
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09-FEB-1996;
                                                                                     diagnosing this potential, the responsiveness of interferon treatment may
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96US-00599252.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kimmel BE,
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밁 Query Match
Best Local Similarity
Matches 19; Conserv 935 21 CTCTATTGCCCAGGCTGGAGT CTCTGTTACCCAGGCTGGAGT 955 Conservative 1.8%; 0; Mismatches Score 17.8; Pred. No. 1. H 1.5e+03 DB 1; Length Indels <u>,</u> Gaps 0

Sequence

22 BP; 6 A; 7 C; 6 G; 3 T; 0 U; 0 Other;

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AAT72014 standard;
 DNA;
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HH; mutation; HH-associated al. HHP-19; HHP-29; microsatellite Primer; polymerase chain reaction; amplify; hereditary haemochromatosis; HH-associated allele; base-pair polymorphism; HHP-1; 9; microsatellite repeat allele; genetic marker; eatment; hepatitis C infection; ss.

Primer detects marker 950-8 in HH region of chromosome

6p2.1.

18-AUG-1997

(first entry)

ARESULT 1006
AAT72014/c
ID AAT7201
XX AAT7201
XC AAT7201
XC AAT7201
XC Primer
XX Primer
XX Primer
XX Primer
XX HH; mut
XW HH; mut
XW HHP-19;
XW interfe
XX Synthet
XX PN WO96356
XX
PN WO96356
XX
PF 06-MAYXX 06-MAY-1996; Synthetic. 14-NOV-1996. 96WO-US006352

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The sequences given in AAT71973-2044 represent a series of primer pairs CC which were used to determine the presence or absence of the common CC hereditary haemochromatosis (HH) gene mutation in an individual The CC method comprises assessing genomic DNA from an individual for the CC method comprises assessing genomic DNA from an individual for the CC presence or absence of the HH-associated allele of the base-pair CC polymorphism HHP-1, HHP-19 or HHP-29, and/or at least one non-optional CC marker comprising the following microsatellite repeat alleles of group A CC and optionally of group B: Group A: 19D9(205), 18B4(235), 1A2(239), CC 1E4(271), 24E2(245), 2B8(206), 3321-1(197), 4073-1(182), 4440-1(180), CC 1(142), 950-2(164), 950-3(165), 950-4(128), 950-5(180), 950-6(151), 950-4(128), 950-1(148), 950-6(151), 950-4(128), 950-5(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-1(180), 65-
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                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 1007
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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15-NOV-1995;
09-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Determn. of the common hereditary haemochromatosis gene primers based on novel microsatellite repeat flanking se base-pair polymorphisms HHP-1, HHP-19 or HHP-29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 14; Page 15; 67pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MERC-) MERCATOR GENETICS INC
                                                                                                                                                               polymorphism; biallelic; human; forensic; paternity testing; disease; detection; phenotypic typing; characteristic; infection; hereditary; autoimmune disease; cancer; inflammation; drug; therapy; medicament; treatment; marker; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                            AAX09910 standard; DNA; 22 BP
                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                   Human biallelic polymorphic marker downstream primer #216.
                                                              WO9820165-A2
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19; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTCACTCTGTTACCCAGGCTG 951
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                                                                                                                                                                                                                                                                                                            (first entry)
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95US-00559302.
96US-00599252.
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Pred. No. 1.5e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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PAXX PRAXX P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC genome (represented in AAXI0269-X12937). These primers can be used in a comethod for determining polymorphic forms in an individual for use in e.g. CC method for determinity testing or phenotypic typing for diseases such CC as agammagiobulinemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular CC dystrophy, Wiskott-Aldrich syndrome, Fabry's disease, familial CC hypercholesterolemia, polycystic kidney disease, hereditary CC sphercytosis, von Willebrand's disease, tuberous sclerosis, hereditary CC haemorrhagic telangiectasia, familial colonic polyposis, Ehlers-Danlos CC syndrome, osteogenesis imperfecta, acute intermittent porphyria, CC autoimmune diseases, inflammation, cancer, diseases of the nervous classification by pathogenic microorganisms, and characteristics such cas longevity, appearance (e.g. baldness, obesity), strength, speed, CC endurance, fertility, and susceptibility or receptivity to particular CC drugs or therapeutic treatments. The isolated polymorphic nucleic acid commonates can also be used to produce medicaments for the treatment or produce medicaments for the treatment or produce medicaments or the treatment or produce medicaments for the treatment or produce medicaments.
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AAX89393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX09121-X10268 are allele-specific oligonucleotide primers used in the isolation of various biallelic polymorphic markers found in the human genome (represented in AAX10269-X12937). These primers can be used in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6-NOV-1996;
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                                                         20-JAN-1998;
09-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                         mammary associated chemokine; MACK; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chemokine; breast tissue; breast milk; breast disease; vaccine; human; inflammation; infection; mastitis; benign cystitis; hyperplasia;
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                                                                                                                                                  12-JAN-1999;
                                                                                                                                                                                                              22-JUL-1999
                                                                                                                                                                                                                                                                          WO9936540-A1
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19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                         98US-0071899P.
98US-0092155P.
                                                                                                                                                        99WO-US000651.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
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Pred. No. 1
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(CODO-) CODON

DIAGNOSTICS LLC

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RESULT 1009
AAH40206(c)
ID AAH4020
XX AAH4020
XX AAH4020
XX ID Spe
XX Single
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Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A mammary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Papsidero
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 28; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; tesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney dieease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                  New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Single nucleotide polymorphism; SNP; single nucleotide primer extension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP specific lower PCR primer SEQ ID 3002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH40206 standard;
                                                                                                                                                                                     Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                            15-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inflammation;
                                                                                                                                                                                                                                                    (ORCH-) ORCHID BIOSCIENCES INC
                                                                                                                             2001-290930/30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  forensic
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                                                                                                                                                                                                                                                                                                            99US-0160096P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               investigation; paternity analysis; PCR primer; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                        nucleic
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Crimeric Annivers and the sequences of regions flanking concludes kits for determining the presence or absence of a SNP, using the presence or absence of a SNP, using the coligonucleotides of the invention. The PCR primers are used to amplify a CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or CC oligonucleotides are useful for determining the presence, absence or CC oligonucleotides are useful for determining the presence, absence or CC oligonucleotides are useful for determining the presence, absence or CC oligonucleotides are useful for determining the presence, absence or CC oligonucleotides apathological phenotypic trait suspected of being CC caused by one or more SNPs. Phenotypic traits include diseases e.g. CC agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular CC diseases including, diabetes insipidus, Lesch-Nyhan syndrome, muscular CC osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial CC diseases, including, rheumatoid arthritis, multiple sclerosis, cc inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and contermity analysis. The present sequence represents a PCR primer specific for a human SNP containing NNA sequence.
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 65; 83pp; English.
                                                                                                                     human SNP
22 BP; 7
                                                                                                              containing
A; 6 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAH40944 represent PCR primers, single nucleotide
                                                                                                                     DNA sequence
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RESULT 1010 AAD31451 Query Match Best Local S Matches 19 AAD31451; AAD31451 standard; DNA; 22 BP 695 21 19; Similarity CAGGTTCAAGTGATTCTCCTG CGGGTTCAAGTTATTCTCCTG 715 Conservative 1.8%; 0 Score 17.8; Pred. No. 1 ed. No. 1.5 1.5e+03; BB 1: 2 Length Indels 0, Gaps

0

밁 5

31-MAY-2002 (first entry)

Human chromosome 17 92Kb gene fragment amplifying PCR primer, Span2F.

Human; Van Buchem's disease; genomic deletion; craniotubular hypertosis; autosomal recessive disorder; chromosome 17; chromosome 17q21; bone dysplasia; 92Kb gene fragment; PCR primer; ss.

WO200210455-A2

07-FEB-2002.

30-JUL-2001; 2001WO-US023968

06-JUL-2001; 28-JUL-2000; 2000US-0221855P 2001US-0303386P

CELLTECH R STRAEHLING HAMPTON K. & D INC

Ħ, Prol1 S Paeper В;

WPI; 2002-227089/28.

Methods for identifying subjects who are afflicted with or carriers of diseases associated with genomic deletion(s), e.g. Van Buchem's disease by determining the presence of a deletion in the 92 kb region of human disease, f human

sample.

The present individuals

invention relates to methods for distinguishing between homozygous for and therefore afflicted with Van Buchem's

homozygous for and therefore

Claim chromosome

Page 26; 109pp;

English

17

at

disease, individuals heterozygous for and therefore carriers of Van Buchems's disease and individuals who are not afflicted with Van Buchems's disease comprise identifying a large genomic deletion in chromosome 17 at 17g21. The method is useful for identifying individuals who are afflicted with or carriers of diseases associated with one or more genomic deletion, particularly Van Buchem's disease, which is a rare autosomal recessive disorder that results in a bone dysplasia referred to a cranictubular hypertosis. The present sequence is a PCR primer used to amplify 92Kb gene fragment in human chromosome 17 at 17g21

Matches Query Match

l Similarity 19; Conserv

Conservative

0;

1.8%;

Score 17.8; pred. No. 1 Mismatches

.5e+03; DB 1; 2

Length Indels

0

Gaps

0

문 S 22

B₽;

5 A;

10 C; 3 G; 4 T; 0 U; 0 Other;

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RESULT 1011
ABK65937/c
ID ABK6593
XX ABK6593
XX ABK6593
XX DT 02-JUL-
XX Primer;
XX Prim
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The invention relates to producing a sub-population of labeled nucleic acids (NAs) comprising contacting a NA sample from a physiological source, with a pool of 50 distinct gene specific primers under suitable conditions to enzymatically generate sub-population of NAs, where each gene specific primer has a sequence complementary to a distinct mRNA, and each labeled NA is generated using a single gene specific primer. The method is useful for producing a sub-population of labeled NAs which is useful for analysing the differences in the RNA profiles between several different physiological sources, where the method comprises producing subpopulation of labeled NAs for the different physiological sources, comprising the populations for each physiological source to identify differences in the population, where the comparison is preferably
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Producing sub-population of labeled nucleic acids, useful for analyzing differences in RNA profiles between several different physiological sources, using set of distinct gene specific primers.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human gene specific PCR primer #25
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                                                   performed by hybridising the labeled NAs for each of the distinct physiological sources to an array of probe NAs stably associated with the surface of a substrate to produce a hybridisation pattern for each of the sources, and comparing the patterns for each of the sources, where differential gene expression assays are utilised in differential expression analysis of diseased a normal tissue e.g. neoplastic a normal tissue, or different tissue or subtissue types. The present sequence is a human gene specific PCR primer used in the method of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from USPTO at http.wipo.seqdata.uspto.gov/sequence.html?DocID=6352829B1
A; 10 C; 4 G; 3 T; 0 U;
            0 Other;
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Matches Query Match Best Local 643 CCCAGGCTGGAGTGCAGTGGC 663 19; 21 Similarity Conservative 1.8%; 90.5%; 0; Score 17.8; Pred. No. 1. Mismatches .5e+03; DB 1; Length Indels 0 Gaps

0

Sequence

22 BP; 5

AAD43557 AAD43557; Human CD2000 DNA amplifying 14-NOV-2002 standard; DNA; 22 (first entry) forward primer.

RASULT 1012
AAD43557/c
ID AAD4355
XX Human C
XX Human C
XX Human;
KW immune
KW carcinc
KW osteopas
KW dabbete
KW emphyse
KW emphyse
KW immunos
KW jaundic
XX
PR O3-NOVPR 02-NOVXX
PR 03-NOVXX
PR 03-Human; immunoglobulin; Ig; SLAM associated protein; SAP; CD2000 protein; immune proliferative disorder; immune disorder; rheumatoid arthritis; carcinoma; autoimmune disorder; multiple sclerosis; Grave's disease; Hashimoto's disease; acquired immune deficiency syndrome; hepatotropic; osteoarthritis; allergic inflammatory disorder; viral infection; asthma; psoriasis; apoptotic disorder; systemic lupus erythematosus; bronchitis; diabetes mellitus; septic shock; chronic obstructive pulmonary disease; emphysema; cachexia; hepatic circulatory disorder; hepatitis; cirrhosis; acute myeloid leukaemia; haemophilia; anaemia; gene therapy; cytostatic; immunosuppressive; neuroprotective; antiinflammatory; Crohn's disease; osteopathic; antibacterial; immunosuplatic; inflammatory bowel disease; osteopathic; antibacterial; immunosuplatic; inflammatory bowel disease; jaundice; dermatological; ulcerative colitis; AIDS; PCR; primer;

EP1223218-A1 sapiens

17-JUL-2002.

02-NOV-2001; 2001EP-00309339

03-NOV-2000; 2000US-00706167

Fraser

(MILL-)

MILLENNIUM PHARM INC

2002-620680/67.

Novel isolated polypeptide containing immunoglobulin and immunoglobulin-like domains and SLAM associated protein, termed CD2000 or CD2001, usefu for treating immune, inflammatory, or hepatic circulatory disorders. useful

Disclosure; Page 75; 138pp; English.

The invention relates to nucleic acid moleculencedes a polypeptide containing immunoglobuland SLAM associated protein (SAP) motifies. Cluseful for treating disorder such as immune immune disorders (e.g. carcinoma), viral inf acid molecule, immunoglobulin acid molecule, designated CD2000 wh immunoglobulin (Ig) and Ig-like dom motifs CD2000 DNA and protein is as immune proliferative disorders, viral infection, autoimmune disor. disorders domains which

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autoimmune disease

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RESULT 1013
AAD63370/c
ID AAD6337
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; CD2000; CD2001; therapy; THI disorder; insulin-dependent diabetes; chronic inflammatory disease; organ specific autoimmunity; sarcoidosis; graft rejection; lymphoproliferative disorder; psoriasis; leukaemia; immune disorder; graft versus host disease; inflammatory bowel disease; contact dermatitis; Chron's disease; ulcerative colitis; infection; autoimmune disease; multiple sclerosis; inflammatory disorder; asthma; rheumatoid arthritis; chronic obstructive pulmonary disorder; bronchitis; cystic fibrosis; bronchiolitis; hypersensitivity pneumonitis; emphysema; lung cancer; idiopathic pulmonary fibrosis; pneumonia; hepatic failure; jaundice; hereditary hyper bilirubinaemia; hepatic circulatory disorder; hepatitis; malignant tumour; hepatic vein thrombosis; colon cancer; amyloidosis; cirrhosis; lymphoma; scleroderma; mastocytosis; anaemia; heamyloidosis; thalassaemia; dermatological; cytostatic; neuroprotective; immunosumposasive. hepatityping. PCP: neuroprotective; immunosumposasive. hepatityping.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAD63370 standard; DNA; 22
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                                                                                                                                                                                                                              03-NOV-2000; 2000US-00706167
02-NOV-2001; 2001US-00007303
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                                                                                                                                                    (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                      12-MAY-2003; 2003US-00436523
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ımmunosuppressive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1005 CGATTCTCCTGTCTCAGCCTC 1025
2003-843934/78
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hepatotropic; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       В₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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A new nucleic acid designated CD2000 encodes a polypeptide containing and Ig-like domains and a SLAM associated motif and is useful to treat TH1 disorders including chronic inflammatory disease, diabetes and

Ιg

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RESULT 1014
ADI23730
   PIX PAR RECENT OF THE PROPERTY OF THE PROPERTY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to novel CD2000 and CD2001 proteins and CD2001 proteins and CD2001 proteins and polynucleotides encoding such proteins. Sequences of the invention are CC used to treat TH1 disorders, particularly chronic inflammatory diseases, insulin-dependent diabetes, organ specific autoimmunity, psoriasis, graft rejection, contact dermatitis, graft versus host disease or sarcoidosis. The invention is useful to modulate or to identify modulators of immune CC disorders such as lymphoproliferative disorders (e.g., leukaemia or and X -linked lymphoproliferative disease), inflammatory bowel disease such as chronic disease and ulcerative colitis, autoimmune disease such as CC multiple selerosis, inflammatory disorders such as rheumatoid arthritis and asthma, chronic obstructive pulmonary disorders and viral, bacterial, fungal or parasitic infections. The invention is also useful to identify, CC isolate, deplete, track, or modulate the differentiation, replication cCC analso be used to modulate the function, morphology, proliferation also be used to modulate the function morphology, proliferation cCC and therefore can be used to treat disorders such as bronchitis, cystic fibrosis, bronchiolitis, hypersensitivity pneumonitis, emplysema, lung cancer, idiopathic pulmonary fibrosis, pneumonia, jaundice, hepatic failure, hereditary hyper blirubinaemias, hepatic circulatory disorders, malieronari rumours, henatic vair rhrombonia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                      (WENX/)
(STEW/)
(TSUI/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      lipase; LPDLR; LPDLR; lipase deficiency; atherosclerosis; fatty liver disease; dyslipidaemia; hypercholesterolaemia; hypertriglyceridaemia; mixed dyslipidaemia; lipid deficient lipoprotein deficient state; human; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human LPDLR PCR primer #10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADI23730 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hepatitis, cirrhosis, malignant tumours, hepatic vein thrombosis, lymphoma, leukemia, colon cancer, amyloidosis, scleroderma, mastocytosis, haemophilia, anaemia and thalassaemias. The present sequence is human CD2000 cDNA specific PCR primer used in the invention
Wen X,
                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003055995-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                              21-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                        10-JUL-2003
                                                                                                                                                                                                                 10-JAN-2002;
                                                                                                                                                                                                                                                                                                           23-DEC-2002; 2002WO-CA001998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1005 CGATTCTCCTGTCTCAGCCTC 1025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19;
                                                   TSUI L.
HEGELE R A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                    WEN X
Stewart AK,
                                                                                      STEWART A K.
TSUI L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CGATTCTCCTGCCTCAGTCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                 2001US-0341786P
2002US-0346603P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; 3 C; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90.5%;
Tsui L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Opp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  G; 2 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 17.8; DB 1;
Pred. No. 1.5e+03;
0; Mismatches 2;
   Hegele RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             state;
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WPI;

2003-569444/53

identifying

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ABV77329/c
II ABV77
XX
AC ABV7
AC ABV7
DT 07-F
XX
Huma
XX
Homo
OS Homo
OS Homo
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FPN CN13
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PD 03-f
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PF 12-f
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CC lipase polypeptide (polyp), e.g., LPDL (I) or LPDLR polyp (II). (I) or
CC (II) is useful for identifying substances which can bind with LPDL or
CC LPDLR polyp, and for identifying a compound that affects the binding of
CC their nucleic acid is useful for identifying a compound that affects lPDL or
CC their nucleic acid is useful for identifying a compound that affects LPDL
CC or LPDLR polyp activity or expression. (I) or (II) or their nucleic acid
CC increased or detecting or monitoring a condition associated with
CC increased or decreased LPDL or LPDLR expression or activity in an animal,
CC where the condition is lipase deficiency, atherosclerosis, fatty liver
CC disease and dyslipidemias, such as hypercholesterolemia,
CC hypertriglyceridemia, mixed (combined) dyslipidemia, lipid or lipoprotein
CC deficient states, and/or any other tissue or plasma disorders of lipid or
CC lipoprotein metabolism. The nucleic acid is useful for diagnosing the
CC involves detecting a germline alteration in the nucleic acid in the
CC subject. An inhibitor is useful for modulating triglyceride activity by
CC inhibiting expression or activity of (I) or (II). The nucleic acid is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel isolated LPDL or LPDLR lipase polypeptides, useful for identify substances that bind to the protein and which are useful for treating diseases associated with lipase function e.g. atherosclerosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hypercholesterolemia.
                                                                                                                                                                                                           12-SEP-2000; 2000CN-00125186
                                                                                                                                                                                                                                                                                CN1342770-A
                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                      Human protein
                                                                                                                                                                                                                                                                                                                                                                                                                          07-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABV77329;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABV77329 standard; DNA; 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          exemplification
                                                                                                                                                                          12-SEP-2000;
                  New human protein 10.01 of encoding polynucleotide,
                                                                        WPI; 2002-529811/57.
                                                                                                         Ķ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      as a probe or primer. The present sequence is used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N
                                                                                                                                                                                                                                                                                                                                                     10.01;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                           Xie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AACTCCCGACCTCAGATGATC 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AACTCCTGACCTCAGGTGATC 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₿₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                            2000CN-00125186
                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                                                                                                                                                                     aminolyase active site; arrhythmia; diabetes; probe;
                                                                                                                                                                                                                                                                                                                                                                                        10.01 related probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ID NO 66; 172pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of the invention.
                                                                                                                                             DEV
                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7 C; 4 G; 5 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90.5%;
                                                                                                                                           გ
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0;

Mismatches

Pred.

No. 1.5e+03;

Score 17.8;

DB 1; 2

Length 22; Indels

0,

Gaps

0

0 U; 0 Other;

ΒP

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containing Phe-His, useful for treatir

Phe-His aminolyase active treating arrhythmia and c

e site and diabetes.

TTD

SHANGHAI

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AAT66003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 7; Page 22 (disclosure); 33pp; Chinese.

The invention relates to a human protein designated 10.01, containing the phe-His aminolyase active site. Also disclosed are the encoding polynucleotide, and a method for preparing the polypeptide by DNA recombination. The application of the polypeptide is in treating arrhythmia and diabetes. Also disclosed are the antagonist against polypeptide and its therapeutic action, and the application of the polypeptide. The current sequence represents a human protein 10.
                                  The invention relates to the isolation of polymorphic repeat sequences having the sequence (dC-dA)n. (dG-dT)n which can be used as genetic markers. Primers based on these sequences can be used to detect these repeats, especially for use in e.g paternity or maternity testing, huma genetic analysis such as linkage analysis of genetic disease, commercia animal or plant breeding or pedigree analysis. Clones containing the repeat sequences were isolated by hybridisation of chromosome-specific phage libraries with a synthetic poly(GC-dA). (dG-dT) probe. Over 100 repeat blocks were isolated. The primers ANT65798-766047 were used to Pamplify the inserts from the isolated clones containing the repeat sequences. The primers ANT66002-3 were used to amplify the repeat primers ANT670410 (ANTA) and the repeat primers ANT670410 (ANTA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 41 BP; 6 A; 17 C; 9 G; 9 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT66003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT66003
                     sequence marker
                                                                                                                                                                                                                                                                                                                       Claim 7; Col 13-14; 186pp; English.
                                                                                                                                                                                                                                                                                                                                                                 Detection of polymorphic genetic markers of the using novel nucleic acid mols. as primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1997-042299/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Weber JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-APR-1989;
05-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-DEC-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US5582979-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR; polymerase chain reaction; paternity; maternity; human; pedig
linkage analysis; genetic disease; animal; plant; breeding; locus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Primer #2 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAR-2003
18-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MARS-) MARSHFIELD CLINIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 651 GGAGTGCAGTGGCGCAATCTTGGCTCACTGCA 682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GGGTTGCAGTGGACCAAGATTGCGCCACTGCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amplify repeat sequence marker Mfd103.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               repeat sequence; genetic marker; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       89US-00341562.
91US-00754351.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94US-00222177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chromosome;
                     clone Mfd103 (AAT65774).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.8%;
71.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 17.6;
Pred. No. 1
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                        (Updated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein 10.01
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RESULT 1018
AAA60279/c
ID AAA6027
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AC AAA6027
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AAZ35377/c
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                                                                                                                        Query Match
Best Local Similarity
Matches 18; Conserv
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  AAA60279;
                       AAA60279 standard;
                                                                                                                                                                   Sequence 19
                                                                                                                                                                                                 The present sequence is that of interspersed repeated sequence PCR (IRS-PCR) primer ALU3' used to identify human-specific sequences in yeast artificial chromosomes (YAC) derived from the human chromosome 1923.3-q24 region. The chromosomeal region contains the locus associated with absorptive hypercalciuria (AH). IRS-PCR fingerprints were generated, and genes contained within YACs were identified by exon trapping. cDNA corresponding to the AH gene was isolated (see AAZ33376). Identification of the AH genomic region allows genetic screening for increased risk of
                                                                                                                                                                                                                                                                                                                           Novel genomic region useful in screening osteoporosis with hypercalciuria.
                                                                                                                                                                                                                                                                                                                                                                                  Reed-Gitomer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               W09967426-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        osteopathic; anticalciuric; chromosome PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ35377;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-116959/10.
                                                                                                                                                                                                                                                                                                                                                                                                                             23-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; absorptive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Interspersed repeated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ35377
                                                                                                                                                                                        developing
                                                                                                                                                                                                                                                                                                                                                                                                       (TEXA ) UNIV TEXAS SYSTEM.
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                                                                                                   651
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                                                                                                  GGAGTGCAGTGGCGCAATC 669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
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                                                                              GGAGTGCAGTGGCGCGATC 1
                                                                                                                                                                                                                                                                                                      Page 125; 153pp; English.
                                                                                                                                                                                          AH or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 3 A;
                                                                                                                                                                   BP; 3 A; 9 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                 BΥ,
                                                                                                                                                                                                                                                                                                                                                                                                                             98US-0090348P
                                                                                                                                                                                                                                                                                                                                                                                                                                                   99WO-US014347
                                                                                                                                                                                         osteoporosis with hypercalciuria
                                                                                                                                                                                                                                                                                                                                                                                  Pak
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hypercalciuria; osteoporosis; nephrolithiasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                       DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.8%;
                                                                                                                                  94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8 C; 3
                                                                                                                                                                                                                                                                                                                                                                                  CYC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence PCR
                                                                                                                                               1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19
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                       ΒP
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                                                                                                                                     Pred.
                                                                                                                                               Score
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                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17.4;
No. 1
                                                                                                                                     No.
                                                                                                                                             17.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer ALU3'.
                                                                                                                                  1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1q23.3-q24; therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .4e+03;
                                                                                                                                                                                                                                                                                                                                      for absorptive hypercalciuria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 19;
                                                                                                                                             Length 19;
                                                                                                                         Indels
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                                                                                                                       Gaps
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RESULT 1019
AAA48211/c
ID AAA4821
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         murine prostate cancer predisposing genes HPC2 and Mm.HPC2. The human version of the gene is found on chromosome 17p. Some alleles cause a predisposition to cancer, particularly prostate cancer. This gene and it protein can be used in peptide and gene therapy for cancer patients, as well as being useful as diagnostic tools (both for cancer sufferers and those with a predisposition to the disease) and in the production of
                                                                                                       cardiovascular disease; coronary artery disease; non-insulin dependent diabetes mellitus; neuropathy in NIDDM; essential hypertension; hyperlipidemia; diabetic neuropathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 3 A; 2 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human prostate cancer (HPC)2 nucleic acids, polypeptides, and antibodies, useful for treatment and diagnosis of prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; mouse;
human chromoso
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-DEC-2000
                                                                                                                                               Tumour necrosis factor; TNF; TNF-R2; TNFRSF1B; PCR tumour necrosis factor receptor superfamily member
                                                                                                                                                                                                                    15-SEP-2000
                                                                                                                                                                                                                                               AAA48211;
                                                                                                                                                                                                                                                                         AAA48211 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence is a primer used in the isolation of the human and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 5; Page 59; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tavtigian SV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer; sequencing
                          WO200031293-A1.
                                                                             vasoprotective; antihypertensive;
D1S2834; D1S2728; ss.
                                                                                                                                                                                        Reverse PCR primer
                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MYRI-) MYRIAD GENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000-376481/32
                                                                                                                                                                                                                                                                                                                                                                      541
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; mouse; prostate cancer predisposing gene; HPC2;
chromosome 17p; gene therapy; peptide therapy; drug design;
rimer; sequencing primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HPC2
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                                                                                                                                                                                                                                                                                                                                           CCTCAGCCTCCCAAATAGC
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                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0107468P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 exons 2/3 mutation screening primer SEQ ID NO: 100.
                                                                                                                                                                                           for detection of microsatellite
                                                                                                                                                                                                                                                                                                                                                                                                              94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Simard J,
                                                                                                                                                                                                                                                                                                                                                                                                 <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                      559
                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                         Score 17.4;
                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                            lipid-lowering; chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rommens
                                                                                                                                                                                                                                                                                                                                                                                                               1.4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                primer;
1B; huma
                                                                                                                                                                                           marker
                                                                                                                                                   human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This gene and its
                                                                                                                                                                                            D1S2728.
                                                                                                                                                                                                                                                                                                                                                                                                 <u>,</u>
                                                                                            1p36.2;
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02-JUN-2000.

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RESULT 1020
AAF59729
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A novel method for determining a predisposition in a subject to a complex CC polygenic disease involves assaying chromosome 1 for a genetic marker CC indicative of a predisposition to the disease. This method may be used CC for determining predisposition to the disease. This method may be used CC determining predisposition to cardiovascular disease, coronary artery CC disease, non-insulin dependent diabetes mellitus, neuropathy in NIDDM, CC essential hypertension, hyperlipidemia and diabetic neuropathy. The CC method can be used for testing an individual with a family history or in CC the early stages of a complex polygenic disease to ascertain the chance of developing hypertension, neuropathy or lipid disturbances such as high CC total cholesterol, high low density lipoprotein cholesterol, abnormal CC total cholesterol, high low density lipoprotein cholesterol, abnormal CC apolipoprotein Al and abnormal glycosylated haemoglobin. Once a complex CC polygenic disease disposition has been identified the subject can be trasted to prevent or reduce the disease or delay its onset. The genetic CC marker used in the method is DIS2334 and includes a CA repeat region in CC (INFRSF1B) gene. The marker is located at chromosome 1936.2. The present complex equence is the reverse PCR primer used for detection of the mired to mirror as found to be linked to mirror to be complex primer used for detection of the complex processes and the complex processes and the complex processes and the present complex processes and the present processes and the processes 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       e.g. coronary heart disease, hyperhipheemia and how insurant were diabetes mellitus comprises assaying chromosome 1 for a genetic marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 45; 50pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UNSY ) UNIV SYDNEY
                                                                                                                                                                                                                                                           pulmonary embolism; deep vein thrombosis;
                                                                                                                                                                                                                                                                            Protease-activated receptor 4; PAR4; human; activity modulation; thrombin-mediated platelet activation; inhibitor; antagonist; thrombotic disorder; thromboembolism; myocardial infarction; stroke;
                                                                                                                                                                                                                                                                                                                                                                            Human protease-activated receptor 4 (PAR4) RT primer,
                                                                                                                                                                                                                                                                                                                                                                                                                             27-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF59729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          microsatellite marker D1S2728. This marker was found to be linked to
                           24-AUG-1999;
                                                                       01-FEB-2001
                                                                                                                   WO200107072-A1
                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                       peripheral arterial occlusion; activator; coagulation disorder,
                                                                                                                                                                                                                 reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     coronary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for diagnosing a predisposition to a complex polygenic disease oronary heart disease, hyperlipidemia and non-insulin-dependent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                              transcription; RT primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACCCAGGCTGGAGTGCAGT 660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACCCAGGCTGGAGTGTAGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                           99WO-US019158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 17.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                 SEQ
                                                                                                                                                                                                                                                                                                                                                                                   ID NO:13
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platelet activation. The method comprises the adminstration of specific modulators of protease-activated receptor 1 (PAR1) and protease-activated receptor 2 (PAR4) activity. The invention also encompasses an anti-PAR4 creatibody directed against all or part of a thrombin-binding site of PAR4. The method is useful for reducing the level of a thrombin response in a mammal or for preventing disorders such as thromboembolism in individuals with a history of thrombosis. Inhibitory compositions are useful in the treatment of disorders such as myocardial infarction, stroke, pulmonary cumbolism, deep vein thrombosis (DVT), peripheral arterial occlusion and cother blood system thromboses. Activating compositions are useful in the treatment of disorders involving insufficient clotting, where dual activation of PAR1 and PAR4 may increase activation of platelets, since thrombin has the ability to activate both receptors. A PAR4 antibody is thrombin mediated platelet activation. The present sequence represents to thrombin-mediated platelet activation. The present sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a method of modulating thrombin-mediated platelet activation. The method comprises the adminstration of specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Affecting platelet activation, for treating e.g. thromboembolism or pulmonary embolism, comprises administering two compounds that modulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Coughlin SR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protease-activated receptor 1 and 4 activity, respectively.
                                                invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 15; 46pp; English.
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밁 Ś Matches Query Match Sequence 19 Local 727 TGAGTAGCTGGGACTACAG 745 18; Similarity TGAGTAGCTGGGATTACAG 19 Conservative BP; 5 A; 1.8%; 2 C; 7 Score 17.4; D: Pred. No. 1.4e 0; Mismatches G; 5 0 ; 0 U; 0 Other; 1.4e+03 DB 1; Length 19; Indels 0; Gaps 0

RESULT 1021 AAH38445/c SNP specific upper PCR primer SEQ ID 1241. 14-AUG-2001 AAH38445 standard; DNA; 19 (first entry)

Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; s single nucleotide primer extension;
a; diabetes insipidus; cancer; primer; ss

Homo

WO200129262-A2

26-APR-2001

13-OCT-2000; 2000WO-US028436

15-OCT-1999; 99US-0160096P.

(ORCH-) ORCHID BIOSCIENCES INC

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RESULT 1022
AAH38669
ID AAH3866
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                         Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP specific
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                                                15-OCT-1999;
                                                                                              13-OCT-2000; 2000WO-US028436.
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(ORCH-) ORCHID BIOSCIENCES INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              upper PCR primer SEQ ID 1465.
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                                                99US-0160096P
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CC primer extension (SNPE) primers, and the sequences of regions flanking considers of single nucleotide polymorphisms SNPs. The present invention considers with the for determining the presence or absence of a SNP, using the cligonucleotides of the invention. The PCR primers are used to amplify a component state of ignorphisms of the invention. The PCR primers are used to amplify a component state of the invention. The PCR primer are used to amplify a component state of the invention. The primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The coligonucleotides are useful for determining the presence, absence or considered the primer extension reaction. The constant of a SNP and for genotyping nucleic acid samples, for e.g. to disently of a SNP and for genotyping nucleic acid samples, for e.g. to cases by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic traits include diseases e.g. cagamaglobulinaemia, diabetes insipidus, teach-Nyhan syndrome, muscular constants also include symptoms of or susceptibility to multifactorial constants also include symptoms of or susceptibility to multifactorial confissase of which a component is or may be genetic such as autoimmune confissases, including, rheumatoid acute intermittent porphyria. Phenotypic inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and confissassion of the present sequence represents a PCR primer specific for a human SNP containing NNA sequence.
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Sequence 19 BP; 4 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
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                                                                                                            containing DNA sequence
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                             Matches
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Best Local
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                                    Similarity
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TAGCTGGGATTACAGGCGC
                             Conservative
                                    94.7%;
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19
                                    Score 17.4;
Pred. No. 1.
              749
                             Mismatches
                                    .4e+03;
                                            ВG
                                          Length 19;
                             Indels
                             0
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AAH38226/c 14-AUG-2001 AAH38226 standard; (first entry) DNA; 19

SNP specific lower PCR primer SEQ ID 1022.

Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; SNPE; genotyping; agammaglobulinaemia; qiabetee insignitus, vincor, lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; polycystic kidney disease; rheumatoid arthritis; multiple sclerosis; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; ss. Homo sapiens. 13-OCT-2000; 2000WO-US028436 inflammation; forensic investigation; paternity

15-OCT-1999;

99US-0160096P

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RESULT 1024
AAH38229
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                                                                                                                                                                                                                                                           SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch Nyhan syndrome; muscular dystropby; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                     Single nucleotide SNPE; genotyping;
                                                                                                                                                                                                                                                                                                                                                                                                             SNP specific upper PCR primer SEQ ID 1025.
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  15-OCT-1999;
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                                                                                                                                         WO200129262-A2
                                                                                                                                                                                                                                          inflammation;
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mes 18; Conserv
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                                             2000WO-US028436
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
99US-0160096P
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                                                                                                                                                                                                                                                                                                                                                            polymorphism; SNP; single nucleotide primer extension;
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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking concludes kits for determining the presence or absence of a SNP, using the Oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The Oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid sample by cassess by association analysis the genotype of an individual or group of cassess by association analysis the genotype of an individual or group of casses by association analysis the genotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. cagammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular cystrophy, familial hypercholesterolaemia, polycystic kidney disease, costeogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial diseases, including, rheumatoid arthritis, multiple sclerosis, confidences, including, rheumatoid arthritis, multiple sclerosis, microgenic method is also useful in forensic investigations and confidence in human SNP containing DNA sequence represents a PCR primer specific for a human SNP containing DNA sequence
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                                          No. 1.4e+03;
                                                  DB 1; Length 19;
                                   Indels
                                  0
                                   Gaps
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Sequence 19

BP; 4 A; 4 C;

7

G; 4 T;

0 α,

0

Other;

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AAH38677 standard; DNA; 19
                              AAH38677;
                                 ΒP
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14-AUG-2001

(first entry)

Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; SNP specific acute intermittent porphyria; inflammation; forensic invest upper PCR primer SEQ ID 1473. investigation; paternity analysis; PCR primer; ss

Homo sapiens.

WO200129262-A2

13-OCT-2000; 2000WO-US028436

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AAH38221
ID AAH
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  RESULT 1026
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The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autotimmune diseases, including, rheumatoid arthritis, multiple sclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a PCR primer specific for a human SNP containing DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 57; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-290930/30
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13-OCT-2000; 2000WO-US028436.
                                                                                                                                                              polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; s
                                                                                                                                                                                                                         Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; SNPE, genotyping; agammaglobulinaemia; tiabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia;
                                         26-APR-2001
                                                                                   WO200129262-A2
                                                                                                                                                                                                                                                                                                                 SNP specific upper PCR primer SEQ ID 1017.
                                                                                                                                                                                                                                                                                                                                                                                                                                              AAH38221 standard; DNA; 19 BP
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                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                           14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  393 TGCTGGGATTACAGGCGTG 411
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TGCTGGGATTACAGGCATG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 4 A; 3 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                           (first entry)
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Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 19
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CC performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or CC identity of a SNP and for genotyping nucleic acid samples, for e.g. to CC assess by association analysis the genotype of an individual or group of CC individuals, having a pathological phenotype traits include diseases e.g. CC agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular CC dystrophy, familial hypercholesterolaemia, polycystic kidney disease, CC osteogenesis imperfecta and acute intermittent porphyria. Phenotypic CC traits also include symptoms of or susceptibility to multifactorial CC disease of which a component is or may be genetic such as autoimmune CC diseases, including, rheumatoid arthritis, multiple sclerosis, CC inflammation, cancer, nervous system diseases and infection by pathogenic CC microorganism. The method is also useful in forensic investigations and CC paternity analysis. The present sequence represents a PCR primer specific
                                                                                                                                                                                                                                                                                                                                                                                                                                 primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequences AAH37205 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 55; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acid sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-290930/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-OCT-1999;
                                             for a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORCHID BIOSCIENCES INC
                                                gnp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0160096P.
                                             containing DNA sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH40944 represent PCR primers, single nucleotide
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밁 Ś Matches Query Match Best Local Similarity 393 18; \_ TGCTGGGATTACAGGCATG 19 TGCTGGGATTACAGGCGTG 411 Conservative 1.8%; ٥. Score 17.4; Pred. No. 1 Mismatches .4e+03 DB 1; Length 19 Indels <u>.</u>. Gaps 0

Sequence 19 BP; 4 A; 3 C; 7 G; 5 T; 0 U; 0 Other;

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RESULT 1027
AAS13576
                        AAS13576 standard; DNA; 19
AAS13576;
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17-DEC-2001

(first entry)

(56CA1)

16-FEB-2000; 2000EP-00870022. 10-APR-2000; 2000US-0195777P. Human; VMGLOM; CA repeat; PAC Reverse PCR primer used to isolate CA repeats from PAC 612c19 16-FEB-2001; 2001WO-EP001760 23-AUG-2001 WO200160856-A2 Homo sapiens glomulin; venous malformation glomangioma; PCR 612c19; ss. primer

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Best Local S
Matches 18
                                                                                                                                                       16-FEB-2000;
10-APR-2000;
22-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to the isolation of novel human and mouse VMGLOM polypeptides (long form and short form), and the nucleic acid molecules encoding them. VMGLOMS (also referred to as glomulins) are a subtype of venous malformations (VMS) called glomangiomas. In humans, VMGLOM has been mapped to chromosome 1p21-22. VMGLOMs and the nucleic acids encoding for them are useful as a medicament or for incorporation into a diagnostic kit. Such medicaments are useful for preventing, treating or alleviating disorders with a vascular component, particularly where alteration of vascular smooth muscle cell phenotype is needed, e.g. varicosities, cardiopathies or cardiomyopathies, cerebral disorders and cancer. The nucleic acids are also useful in gene therapy. The present sequence for reverse PCR primer is used to isolate novel CA repeats from PAC 612c19 (56CA1) clone in the methods of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-557643/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19
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                                                                                                                                                                                                                                                                                                                                          Human; VMGLOM; CA repeat; PAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 71; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                preventing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New VMGLOM genes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYLO-) UNIV CATHOLIQUE LOUVAIN
 New VMGLOM genes and polypeptides, useful in gene therapy or for preventing, treating or alleviating disorders with vascular component, e.g. varicosities, cardiopathies, cerebral disorders or cancer.
                                                                                                                                                                                                                                                  23-AUG-2001
                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                       Forward PCR primer used to isolate CA repeats from PAC 606m5 clone.
                                                                                                                                                                                                                                                                                                                                                                                                                       17-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAS13569 standard; DNA; 19
                                                                                                                                                                                                                    16-FEB-2001;
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                                                                 WPI; 2001-557643/62.
                                                                                                                            (AXTO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 varicosities, cardiopathies, cerebral disorders or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         544 CAGCCTCCCAAGTAGCTGG 562
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18; Conserv
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                                                                                                                          VIND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genes and polypeptides, useful in gene therapy or for treating or alleviating disorders with vascular component,
                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                         2000EP-00870022.
2000US-0195777P.
2000EP-00870320.
                                                                                                                                                                                                                    2001WO-EP001760
                                                                                                                            CATHOLIQUE LOUVAIN
                                                                                                                                                                                                                                                                                                                                            glomulin; venous malformation glomangioma; PCR primer; 606m5; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94.78;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. 1.4e+03
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RESULT 1029
AAH24568
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to the isolation of novel human and mouse VMGLOM polypeptides (long form and short form), and the nucleic acid molecules encoding them. VMGLOMS (also referred to as glomulins) are a subtype of venous malformations (VMS) called glomanglomas. In humans, VMGLOM has been mapped to chromosome 1p21-22. VMGLOMS and the nucleic acids encoding for them are useful as a medicament or for incorporation into a diagnostic kit. Such medicaments are useful for preventing, treating or alleviating disorders with a vascular component, particularly where alteration of vascular smooth muscle cell phenotype is needed, e.g. varicosities, cardiopathies or cardiomyopathies, cerebral disorders and cancer. The nucleic acids are also useful in gene therapy. The present sequence for forward PCR primer is used to isolate novel CA repeats from
The invention relates to a method for determining the metastatic potential of cancer cells derived from a subject with cancer. The method comprises introducing a cancer cell sample into the upper chorioallantoic membrane (CAM) of an avian embryo into which an artificially generated air pocket has been created, incubating the embryo for intravasation to occur, and detecting migration of the cancer cells from the upper CAM to the lower CAM. The present sequence was used to selectively amplify human specific Alu repeat sequences, which will be present in the cancer cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 71; 157pp; English
                                                                                                                                                                                                              Determining the metastatic potential of cancer cells and measuring invasion, comprises introducing cancer cells into the upper chorioallantoic membrane (CAM) and detecting cancer cell migration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; Alu; metastatic potential determination; cancer; chorioallantoic membrane; CAM; avian embryo; intravasat
                                                                                                                                                               Example; Col 11; 24pp; English.
                                                                                                                                                                                                                                                                                                                               Ossowski L;
                                                                                                                                                                                                                                                                                                                                                                                                    04-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                      04-AUG-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US6228345-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cell migration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human Alu sequence-specific primer Alu-Antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                               (MOUN ) MOUNT SINAI SCHOOL MEDICINE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            606m5 clone in the methods of the present invention
                                                                                                                                                                                                 upper CAM to the lower CAM.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                    99US-00366840.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   drug screening; PCR primer; ss.
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Pred. No. 1
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                                                                                                                                                                                                                       ne upper
cell migration from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
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RESULT 1030
ABA82197
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Matches 18
Query Match
Best Local Similarity
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                                                                            Sequence 19
                                                                                                                                                                                             The present invention describes the human Zmax1 gene and the high bone mass (HBM) gene, which are found on chromosome 11q13.3. The Zmax1 and HBM genes have osteopathic activities. The genes can be used in gene therapy, antisense therapy and in the production of vaccines. They can be used in the diagnosis and treatment of bone disorders including osteoporosis, the diagnosis and treatment of bone disorders including osteoporosis,
                                                                                                                                                                                                                                                                                                                                                                                                       New high bone mass (HBM) modulating bone mass for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-APR-2000; 2000US-00543771.
05-APR-2000; 2000US-00544398:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; high bone mass; HBM gene; Zmax1 gene; chromosome 11; 11q13.3; sequence tagged site; STS; osteoporosis; osteopathic; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABA82197;
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                                                                                                                      Paget's disease, sclerostosis, osteomalacia and fibrous dysplasia. ABA82038 to ABA82700 and AAG68168 to AAG68193 represent sequences the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-JUN-2000; 2000WO-US016951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antisense therapy; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zmax1 gene region physical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in metastatic potential
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sclerostosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          640
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ب
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENOME THERAPEUTICS CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TCGCCCAGGCTGGAGTGCA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TCACCCAGGCTGGAGTGCA 658
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 3 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                         BP; 3 A; 2 C; 8 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Little
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           rapy; vaccine; bone disorder; Paget's disease; adapter;
osteomalacia; fibrous dysplasia; PCR primer; linker; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.8%;
1.8%;
94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                          and
the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Recker RR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      map preparation STS marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                          Zmax1 gent
  Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 17.4; DB 1;
Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                       ment of e.g
17.4; DB 1;
No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Johnson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                     e.g. osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                  proteins useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 19;
                       Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         #156.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                    for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                   used
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888888888888888

RESULT 1032 ABL43899/c ID ABL4389 XX '

ÅBL43899

standard;

DNA; 19

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RESULT 1031
AAS99014/c
ID AAS9901
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밁
                                                                Matches
                                                                                Query Match
Best Local
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                                                                                                                                                                            The invention relates to a human prostate cancer predisposing gene coding for an HPC2 polypeptide. The DNA and protein sequences are useful as diagnostic reagents for identifying a mutant HPC2 nucleotide sequence in a suspected mutant HPC2 allele by comparing the sequence of the suspected mutant HPC2 allele with a wild-type HPC2 sequence. The sequences are also useful for detecting an alteration in HPC2, where the alteration is associated with cancer in a human. The method involves analysing an HPC2 gene or an HPC2 gene expression product from a tissue of the human. The HPC2 gene is useful as a marker for prostate cancer and can be used in gene therapy techniques to suppress neoplastic growth of recipient cells which carry the mutant HPC2 allele. The sequences represent primers used in the methods of the invention, cDNA encoding human and mouse HPC2 and contains the methods of the invention, cDNA encoding human and mouse HPC2 and contains the methods of the invention, cDNA encoding human and mouse HPC2 and contains the methods of the invention, cDNA encoding human and mouse HPC2 and contains the methods of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel nucleic acid sequence encoding HPC2 polypeptide, which is marker for prostate cancer, is useful in gene therapy techniques to restore HPC2 normal levels by which neoplastic growth is suppressed in recipient cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; mouse; HPC2; prostate c
gene therapy; prostate cancer
sequencing primer; PCR primer.
                                                                                                                                Sequence 19 BP; 3 A; 2 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                       Example 8; Page 72; 239pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-066599/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-MAY-2000; 2000US-00564805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-MAY-2001; 2001WO-US014602
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human prostate cancer predisposing gene (HPC2) PCR primer #10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-MAR-2002 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAS99014;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAS99014 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pavtigian SV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MYRI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HOSP-)
                                                                                  Local Similarity
                               541
                                                                                                                                                                 encoding HPC2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       392 GIGCIGGGATTACAGGCGI
19
                                                                18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        _
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MYRIAD GENETICS INC. HOSPITAL FOR SICK CHILDREN.
                                 CCTCAGCCTCCCAAGTAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GTGCTGGGATTACAGGTGT 19
 CCTCAGCCTCCCAAATAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HPC2; prostate cancer; neoplastic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Teng DHF, Simard J,
                                                                                                                                                                 paralogues and orthologues
                                                                              94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cancer predisposing
                                                                                                                                                                                                                                                                                                                                                                                                         English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         410
                                 559
                                                                                  Pred. No.
                                                                                                Score 17.4;
                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rommens
                                                                                  1.4e+03;
                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  growth; cyto
; chimpanzee;
                                                                                                  Length 19;
                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gorilla;
                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                0;
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RESULT 1033
ABL44483
ID ABL4448
XX
AC ABL4448
XC ABL4448
XC ABL4448
XC ABL4448
XC Human C
XX
Human C
XX
KW Human;
KW PCR pri
                                                                                                                                                                                                                                                                                                                                                                                                                         밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a method of arraying genome clones. The cmethod comprises: (a) clones of the genomic libraries contained in cmultiwell plates numbered for discrimination are mixed in each of the cmultiwell plates; (b) a primer designed based on the chromosome marker (c) a signal corresponding to the marker is detected from the resultant camplified product to specify the discrimination Nos. of the multiwell confidence is changed so that the same discrimination Nos. of the multiwell confidence is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell confidence; (e) the clones in the multiwell plates of the specified discrimination Nos. to array the multiwell confidence; (e) the clones in the multiwell plates of the specified confidence; (e) the clones in the multiwell plates of the specified conditateral directions; (f) the mixed clones are cultured and the cresultant cultures are amplified by using the above primer; (g) signals are detected from the amplified by using the above primer; (g) signals are detected from the detected result; and (i) the clones are mixed respectively in each wells of longitudinal constituted as the positions on the chromosome and arrayed. The mixed are specified from the detected result; and (i) the clones are constituted as the positions on the chromosome and arrayed. The mixed primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention
                                                                                                                                                                                                                                                                                                      Ş
                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human chromosome 1p36-35 PCR primer SEQ ID NO:943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-APR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABL43899;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 4; Page 23; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Arraying genome clones.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-MAR-2000; 2000JP-00066716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-MAR-2001; 2001JP-00068285
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                                                                                                                                                                        ABL44483 standard; DNA; 19 BP
                                                        Human chromosome 1p36-35 PCR primer SEQ ID NO:1527.
                                                                                                11-APR-2002
                                                                                                                                    ABL44483;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GENO-) GENOTEX YG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RIKA ) RIKAGAKU KENKYUSHO.
  PCR primer; ss.
                      Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                                                      383 CCTCCCAAAGTGCTGGGAT 401
                                                                                                                                                                                                                                                                    19
                                                                                                                                                                                                                                                                                                                                                l Similarity
18; Conserv
                                                                                                                                                                                                                                                                      CCTCCCAAAGTGCTGGAAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 4 A;
                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                         4 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                0.
                                                                                                                                                                                                                                                                                                                                              Score 17.4; DB 1; Length 19; Pred. No. 1.4e+03; O; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                              Gaps
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The present invention describes a method of arraying genome clones. The centrod comprises: (a) clones of the genomic libraries contained in completes numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker centroduct to specify the discrimination Nos. of the multiwell camplified product to specify the discrimination Nos. of the multiwell centroduct to specify the discrimination Nos. of the multiwell centre containing the clones having said marker sequence; (d) the order of the maximum in the specified discrimination Nos. to array the multiwell centre; (e) the clones in the multiwell plates of the specified discrimination Nos. to array the multiwell centre; (e) the clones in the multiwell plates of the specified discrimination Nos. to are mixed respectively in each wells of longitudinal central directions; (f) the mixed clones are cultured and the centre detected from the amplified by using the above primer; (g) signals are detected from the detected result; and (i) the clones are microarray is useful for gene analysis. ABL42957 to ABL45322 represent pCR primers for human chromosome 21922.1, which are specifically claimed for use in the present invention.
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                                                                  Matches
                                                                                  Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JP2001321190-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-MAR-2001; 2001JP-00068285
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                                                                                                                                  Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; Page 34; 528pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Arraying genome clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-MAR-2000; 2000JP-00066716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GENO-) GENOTEX YG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RIKA ) RIKAGAKU KENKYUSHO.
                       194 TCTCCATGTTGGTCAGGCT 212
                                                                   18;
 \vdash
                                                                                     Similarity
 TCACCATGTTGGTCAGGCT 19
                                                                  Conservative
                                                                                                                                     BP; 3 A; 5 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                   1.8%;
                                                                   0;
                                                                                   Score 17.4;
Pred. No. 1.
                                                                     Mismatches
                                                                                   .4e+03;
                                                                                                    DB 1; Length 19
                                                                     Indels
                                                                     0
                                                                     Gaps
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ABL44464/c
ID ABL4446
XX ABL4446
XX ABL4446
XX ABL4446
XX Human C
XX Human C
XX Human C
XX Human S
XX Homo Se
XX PN JP20013
XX PP 20-NOV-
XX PF 12-MAR
                                                                                                              Human chromosome 1p36-35 PCR primer SEQ ID NO:1508
                                                                                                                                                          ABL44464;
                                                                                                                                                                              ABL44464 standard; DNA; 19
                                                                                                                                   11-APR-2002
                                                                                                                                     (first entry)
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12-MAR-2001; 2001JP-00068285

20-NOV-2001 JP2001321190-A. Homo sapiens PCR primer; ss. Human; chromosome

1p36-35; chromosome 21q22.1; genetic analysis; genome;

10-MAR-2000; 2000JP-00066716

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a method of arraying genome clones. The commethod comprises: (a) clones of the genomic libraries contained in method comprises: (a) clones of the genomic libraries contained in committies plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant camplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified condiscrimination Nos. are mixed respectively in each wells of longitudinal and lateral directions; (f) the mixed clones are cultured and the resultant cultures are amplified by using the above primer; (g) signals cand lateral directions; (f) the mixed clones are cultured and the cromation Nos. of the multiwell plates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABIA2957 to ABIA5322 represent PCR primers for human chromosome 136-35 DNA, and ABIA5323 rob ABIA634 creptesent PCR primers for human chromosome 21q22.1, which are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 18; Conserv
                  WPI; 2002-144136/19
                                                                                                                                                                                                               JP2001321190-A
                                                                                                                                                                                                                                                                                                                                      Human chromosome 1p36-35 PCR primer SEQ ID NO:2316.
                                                                                                                                                                                                                                                                                                                                                                                                                                             ABL45272 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; Page 34; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Arraying genome clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-144136/19
                                                                                                                                                                                                                                                                                               Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                                                                                                                           11-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                             ABL45272;
                                                                                                      10-MAR-2000; 2000JP-00066716
                                                                                                                                       12-MAR-2001; 2001JP-00068285
                                                                                                                                                                          20-NOV-2001
                                                   (RIKA ) RIKAGAKU KENKYUSHO (GENO-) GENOTEX YG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GENO-) GENOTEX YG.
                                                                                                                                                                                                                                                                                 primer;
                                                                                                                                                                                                                                              sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTCACTGCAACCTCTGCCT 692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CTCACTGCAACCTGTGCCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 5 A; 3 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 17.4; DB 1;
Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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multiwell plates; (b) a primer designed based on the chromosome marker (c) sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant camplified product to specify the discrimination Nos. of the multiwell amplified product to specify the discrimination Nos. of the multiwell complete containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified condiscrimination Nos. are mixed respectively in each wells of longitudinal conditions; (f) the mixed clones are cultured and the cresultant cultures are amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The mixroarray is useful for gene analysis. ABL42957 to ABL45322 represent PCR primers for human chromosome 21022.1, which are professionally claimed for use in the present invention.
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a method of arraying genome clones. The the decomprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Arraying genome clones.
19 BP; 4
A; 7 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          528pp; Japanese.
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Best Loc Matches Query Match Local 642 ACCCAGGCTGGAGTGCAGT 660 18; Similarity Conservative 1.8%; 0; Score 17.4; DB 1; Pred. No. 1.4e+03; ed. No. 1.4e Mismatches Length 19; Indels 0, Gaps

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19

ACCCAGGCTGGAGTGTAGT 1

ABL59043 RESULT 1036 JP2002095500-A Homo sapiens. Human; allergosis; eosinophil; primer; Nucleotide sequence of a primer. 20-AUG-2002 ABL59043; ABL59043 standard; DNA; 19 BP (first entry

02-APR-2002.

25-SEP-2000; 2000JP-00291316. 25-SEP-2000; 2000JP-00291316

(GENO-) GENOX SOYAKU KENKYUSHO KK. (KOKU-) KOKURITSU SHONI BYOIN INCHO

WPI; 2002-439993/47.

Examining allergosis, involves measuring the expression levels of a specific gene, and comparing it to the levels in the eosinophils of specific gene, a healthy control.

Example 1; Page 14; 20pp; Japanese

The specification describes a method for examining allergosis. The method comprises measuring the expression level of the gene given in ABL59037, and comparing it with the expression level of the gene in the eosinophils

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RESULT 1037
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XX ABK2299
XX O9-APR-
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XX Human Z
XX Human Z
XX Human Z
XX Homo siteopo
KW Island Geno-
KW Osteopo
XX Homo siteopo
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The invention relates to a method for identifying a molecule involved in lipid regulation comprising identifying a molecule that binds to or inhibits binding of a molecule to high bone mass (HBM) or its wild type gene, Zmax1. Compounds identified by the method are useful for treating, Cd dispossing, preventing or screening for normal and abnormal lipid-cassociated conditions, including arteriosclerosis, cardiovascular Cd disease, stroke, and osteoporosis. The compounds may also be used in the treatment or prevention of diabetic atherosclerosis, neurovascular CC conditions caused by plaque build-up, poor circulation due to plaque build-up and associated poor wound healing. The methods may be used in CG gene therapy, pharmaceutical development, and diagnostic assays for bone development disorders. Molecules identified by comparison of Zmax1 and CC development, in diagnosis of human or animal bone disease, and in the treatment of bone diseases. Sequences hak22776-ABK23411 represent cDNA cmolecules encoding human Zmax1 and HBM, and PCR primers, probes, linkers and adapters of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                lipid-associated condition; arteriosclerosis; cardiovascular disease; ss; osteoporosis; atherosclerosis; diabetic atherosclerosis; plaque build-up; neurovascular condition; wound healing; gene therapy; PCR primer; probe; bone development disorder; antiarteriosclerotic; cardiovascular; osteopathic; cerebroprotective.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of a healthy allergosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying molecules involved in lipid regulation, useful diagnosing, treating or preventing e.g., arteriosclerosis, identifying a molecule that binds to high bone mass gene or corresponding wild type gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-APR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-MAY-2000; 2000US-00578900
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABK22994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         y person.
The prese
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 39; 409pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cDNA reverse PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Little RD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CREIGHTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    THERAPEUTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.8%;
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ant sequence represents a primer, which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SCHOOL MEDICINE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer #78.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ٦;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           comprises
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RESULT 1038
ABQ81231/c
ID ABQ8123
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                                    was used to produce a human 14273 probe (see ABQ81233), which was used to examine the expression profile of 14273. It was found that 14273 colored to molecules are expressed at high levels in adipose tissue, e.g. white adipose tissue and brown adipose tissue, as well as in pancreatic islets. They are upregulated during exposure to cold (i.e. under conditions that affect brown or white adipocyte metabolism) and downregulated in genetic models of obseity. The present invention provides 14273 nucleic acids, colypeptides and antibodies useful for the diagnosis and treatment of metabolic disorders including obseity, anorexia, cachexia and diabettes. Also provided are methods for identifying a subject having a metabolic activity, methods for modulating metabolic activity, methods for modulating metabolic activity, methods for modulating metabolic activity, methods for nipolysis in a subject, and a method for modulating lipogenesis or lipolysis in a subject, and a method for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; 14273; metabolic disorder; obesity; diabetes; anorexia; cachexia; anorectic; antidiabetic; anabolic; transgenic animal; gene therapy; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                           Gimeno
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200267868-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
Sequence 19
                                                                                                                                                                                                                                        (see
                                                                                                                                                                                                                                                                              Example 1; Page 61;
                                                                                                                                                                                                                                                                                                                       Identifying a nucleic acid associated with a metabolic disorder, usefu for diagnosing metabolic disorders, e.g. obesity, comprises contacting the sample with a probe comprising at least 25 contiguous nucleotides.
                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-698629/75
                                                                                                                                                                                                                                                                                                                                                                                                                                     (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-FEB-2001; 2001US-0271655P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-FEB-2002; 2002WO-US006131
                                                                                                                                                                                                                                                                                                             sample with a probe 14273 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 392 GTGCTGGGATTACAGGCGT
                                                                                                                                                                                                                                        ABQ81226),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14273
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP;
                             endogenous glucose
                                                                                                                                                                                                                                     sequence is that of forward PCR primer h14273 for human 14273 26), a nucleic acid associated with metabolic disorders. PCR
 BP;
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                                                                                                                                                                                                                                                                              95pp;
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 C; 9
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                                                                                                                                                                                                                                                                                 English.
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  G; 3 T; 0
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Query Match Best Local S Matches 18

Similarity

1.8%; 94.7%;

Score 17.4; Pred. No. 1.

1.4e+03

DB 1;

Length 19;

Indels

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Gaps

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18;

Conservative

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Mismatches

Page

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RESULT 1039
ADH47845/c
ID ADH4784
XX ADH4784
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Vernet CAM, Li L, Landar Ü, Macdougall J, Malyankar Ü, Mi
Tchernev V, Zerhusen BD, Mi
Baumgartner J, Herrmann J,
Baumgartner J, Gerlach V, Grof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-JAN-2001; 2001US-0261376P.
18-JAN-2001; 2001US-0262454P.
18-JAN-2001; 2001US-0262587P.
31-JAN-2001; 2001US-026530P.
14-FEB-2001; 2001US-026595P.
28-FEB-2001; 2001US-02772409P.
16-MAR-2001; 2001US-0276777P.
17-MAY-2001; 2001US-027677P.
27-SEP-2001; 2001US-035306P.
18-OCT-2001; 2001US-0330336P.
09-NOV-2001; 2001US-0345202P.
                      The present invention relates to novel proteins (I) referred to as NOVX, where X is any number from 1 to 18, and their coding sequences (II) (see ADH47704-ADH47759). The proteins and their coding sequences are useful in the manufacture of a medicament for treating a syndrome associated with a human disease, preferably a NOVX-associated disorder such as metabolic disorders, diabetes, obesity, infectious diseases (viral, bacterial, funcial halistics, obesity, infectious diseases (viral, bacterial,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antidiabetic; anorectic; cardiant; hypotensive; antiarteriosclerotic; anorectic; virucide; antibacterial; fungicide; protozoacide; nootropic; neuroprotective; antiparkineonian; anticonvulsant; osteopathic; antiarthritic; antiinflammatory; dermatological; antiasthmatic; antiinflammatory; dermatological; antiasthmatic; antilipaemic; Gene therapy; human; metabolic disorder; diabetes; obesity; viral infection; bacterial infection; fungal infection; viral infection; protozoal infection; anorexia; cancer; cardiovascular disease; neurodegenerative disorder; Alzheimer's disease; parkinson's disease; pentegos; immune disorder; haemacopojetic disorder; inflammatory skin disorder; asthma; dyslipidaemia; NOV14; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADH47845
                                                                                                                                                                                                                                                                                                                                         preventing, dia osteoarthritis,
                                                                                                                                                                                                                                                Example 3;
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                                                                                                                                                                                                                                                                                                                                                                  isolated NOVX polypeptides and venting, diagnosing or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                              2002-698671/75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sapiens
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                                                                                                                                                                                                                                             Page 346; 380pp; English
                                                                                                                                                                                                                                                                                                                                diagnosing or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alsobrook JP, Colman SD, Spytek KA, Boldog F;
Li L, Shenoy S, Casman S, Guo X, Edinger S;
Malyankar U, Patturajan M, Shimkets RA, Pena
Zerhusen BD, Millett I, Miller C, Lepley DM, S
J, Herrmann J, Peyman JA, Gorman L, Mezes P, K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                  polynucleotides, useful for NOVX-associated disorders e
                                                                                                                                                                                                                                                                                                                                         Parkinson's disease
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Rothenberg
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Kekuda R;
henberg M;
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protozoal),

in bone development,

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RESULT 1040
ACC45577
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Best Local S
Matches 18
       The invention relates to novel transgenic animals expressing the high bone mass (HBM) gene, expressing the corresponding wild type HBM gene, comprising an alteration of the gene encoding LRP5 or LRP6, or expressing an LRP5 that is modulated by an altered gene control sequence introduced by homologous or non-homologous recombination. The transgenic animals are for the study of bone density modulation or bone mass modulation. The invention has osteopathic and cytostatic activity. The polynucleotides of the invention may have a use in gene therapy. The transgenic animals and bone mass is modulated relative to non-transgenic animals of the same species in more than one parameter selected from bone density, bone strength, trabecular number, bone size, or bone tissue connectivity. The transgenic animals, nucleic acids and methods are useful for identifying.
                                                                                                                                                                                                                                                                                                                                                                                                                   11-MAY-2001;
17-MAY-2001;
01-FEB-2002;
04-MAR-2002;
                                                                                                                                                                                                                                                        New transgenic animals (e.g. mice), useful as models density modulation, developing drugs for treating or diseases (e.g. osteoporosis), or diagnosing diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; high bone mass; HBM; LRP5; LRP6; transgenic; bone mass modulation gene therapy; bone density modulation; bone strength; trabecular number; bone size; bone tissue connectivity; bone disease; osteoporosis; PCR; osteomalacia; rickets; Paget's disease; neoplasm of the bone; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diseases (hypertension, atherosclerosis), Alzheimer's disease, Parkinson's disease,
                                                                                                                                                                                                               Disclosure; Page 55; 603pp; English.
                                                                                                                                                                                                                                                      diseases (e.g. osteoporosis),
                                                                                                                                                                                                                                                                                                                 WPI; 2003-129278/12
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                                                                                                                                                                                                                                                                                                                                                                        (GENO-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (osteoarthritis), haematopoietic disorders, inflammatory asthma, and various dyslipidaemias. The present sequence for a NOVX sequence.
                                                                                                                                                                                                                                            reduced bone density.
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                                                                                                                                                                                                                                                                                                                                                                        GENOME WYETH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SIS
                                                                                                                                                                                                                                                                                                                                                                                                                  ; 2001US-0290071P.
; 2001US-0291311P.
; 2002US-0353058P.
; 2002US-0361293P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 17.4;
Pred. No. 1.
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                                                                                                                                                                                                                                                        for studying bone preventing bone characterized by
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PR 17-
PR 01-
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Matches 18
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                                                                                                                                                                  The present invention relates to High Bone Mass (HBM), LRP5 (Zmax1) and LRP6 (mutants, which results in a HBM-like phenotype when expressed in a cell. The HBM-like phenotype results in bone mass modulation and/or lipid level modulation. The invention is useful for diagnosing a HBM-like phenotype in a subject and for preparing a composition for modulating bone mass and/or lipid levels in a subject suffering from e.g. osteoporosis. The present sequence is a Sequence Tagged Site (STS) maker, which was used to prepare a physical map of the Zmax1 (LRP5) gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-MAY-2001; 2001US-0290071P.
17-MAY-2001; 2001US-0291311P.
01-FEB-2002; 2002US-0353058P.
04-MAR-2002; 2002US-0361293P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acid comprising a mutation in LRP5 or LRP6, useful diagnosing a HBM-like phenotype in a subject and for preparing composition for modulating bone mass and/or lipid levels in a suffering from e.g. osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-129214/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Allen
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bone mass mo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence tagged site #156 used to prepare Zmax1 (LRP5) gene region
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                                                                                                 Sequence
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18; Conservat
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                                                                                                      19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ic; Gene therapy; High Bone Mass; HBM; LRP5; Zmax1; LRP6; modulation; osteoporosis; STS; sequence tagged site; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anisowicz A,
                                                                                                                                                                                                                                                                                                                                                                                                    Page 62; 629pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 3 A;
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  Conservative
                                                                                                   BP; 3
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                                                                                                   2 C; 8
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Score 17.4;
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                                                                                                   G; 6 T; 0 U; 0 Other;
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Pred. No. 1.
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                                               Length
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RESULT 1043 ADO14391/c ID ADO1439 XX

ADO14391 standard; RNA;

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RESULT 1042
ADL25097
ID ADL2509
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XX INTEST!
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                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    **
                                                                                                                                                                                                                                                                  disease, multiple sclerosis, allergy, asthma and diabetic mellitus), disease, multiple sclerosis, allergy, asthma and diabetic mellitus), and sequence of the immune system, hypersensitivity, analysis, and blood group incompatibility. The present DNA sequence represents a PCR primer that was used to amplify an intestinal epithelium/peyer's patch M cell-associated DNA sequence of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel isolated or purified polypeptide encoded by genes associated with intestinal epithelium or M cell development, differentiation or function, useful for treating autoimmune diseases and infectious diseases.
                                                                                                                                                                                                                                                                                                                                                                                                        invention are also useful in the treatment of: inflammatory bowel disease, glutenenteropathy, infectious diseases, autoimmune diseases (e.g. haemolytic anaemia, rheumatoid arthritis, dermatitis, Grave's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-APR-2001; 2001US-0281416P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention comprises DNA sequences which are associated with intestinal epithelium and peyer's patch M cells. The DNA sequences of thinvention are useful for assessing, modifying, modulating or regulating intestinal epithelium or M cell development. The DNA sequences of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 607; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-075470/07.
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                                                                                                                                                                                                                    Sequence 19
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                                                                                                                                     Similarity
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                                                    CAGGCTGGAGTGCAGTGGC 663
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                                                                                                                                                                                                                       BP; 4 A; 4 C; 8 G;
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                                                                                                                                                                                                                                                                                       invention
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ADO14391;

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                                                                           Matches
                                                                                         Query Match
Best Local
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11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-040878P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cytostatic; vasotropic; nephrotropic; cancer; restenosis; polycystic kidney disease; RNA interference; short interfering RNA; siRNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human interleukin-2-targeted siNA upper strand
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                                                                                                                                                  Sequence 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                      interleukin-2 transcript target sequence.
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19
                                                                         18;
                                                                                           Similarity
                                 GTTACCCAGGCTGGAGTGC 957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    interfering nucleic acid, useful e.g. for treatment and of cancer, downregulates expression of an interleukin gene
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                                                                                                                                                BP; 4 A; 8 C; 5 G; 0 T; 2 U; 0 Other;
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                                                                                         Score 17.4;
Pred. No. 1.
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                                                                                                           DB 1; Length 19;
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XX Cytosta
XW Cytosta
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                                                      The invention relates to short interfering nucleic acids (siNA) which CC downregulate expression of the human interleukin-2 gene by RNA CC interference. The siNAs may or may not comprise ribonucleotides and may CC be double or single stranded. They further comprise sense and antisense CC regions, or alternatively are assembled from a sense oligonucleotide and CC an antisense oligonucleotide. Specifically, the siNAs include short CC interfering RNA (siRNA, double-stranded RNA, micro-RNA (miRNA) and short CC hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, CC can contain deoxyribonucleotides, and can be chemically synthesised, CC expressed from a vector or enzymatically synthesised. The invention also CC relates to kits for the in vitro or in vivo delivery of siRNA; conjugates and/or complexes of siRNA; and vectors that express siNA. The siNAs are CC used to modulate expression of the interleukin-2 gene in cells, tissue CC explants or organisms (e.g., by ex vivo gene therapy), or in grafts and Ct transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis and polycystic kidney disease. The CC identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide CC polymorphisms). The present sequence represents the lower strand of a human interleukin-2-targeted double-stranded siNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of an interleukin gene.
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29-AUG-2002;
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11-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-FEB-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003070744-A1
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2002US-0363124P.

2002US-0386782P.

2002US-0406784P.

2002US-0408378P.

2002US-0409293P.

2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  254; 138pp; English.
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Query Match Best Local Similarity

1.8%; 5 C; 8

Score 17.4; DB 1; Pred. No. 1.4e+03;

Length 19;

Sequence

19

BP;

N ð

G; 0 T; 4 U; 0 Other;

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cc interference. The siNAs may or may not comprise ribonucleotides and may comparison of the human interleukin-2 gene by RNA cc be double or single stranded. They further comprise sense and antisense cc regions, or alternatively are assembled from a sense oligonucleotide and cc an antisense oligonucleotide. Specifically, the siNAs include short cc interfering RNA (siRNA, double-stranded RNA, micro-RNA (miRNA) and short cc interfering respectively. The siNAs can be unmodified or chemically modified, can contain decoxyribonucleotides, and can be chemically synthesised. Cc expressed from a vector or enzymatically synthesised. The invention also cc relates to kits for the in vitro or in vivo delivery of siRNA; conjugates can deduce the conjugates of siRNA, and vectors that express siNA. The siNAs are cc used to modulate expression of the interleukin-2 gene in cells, tissue cc explants or organisms (e.g., by ex vivo gene therapy), or in grafts and ct ransplants for the treatment of a variety of conditions. They may be complicated the construction and contained construction, genetic engineering, therapeutic target cc identification and validation, genetic engineering, pharmacogenomics, cc studying gene function, and gene mapping (e.g., of single nucleotide cc human interleukin-2-targeted double-stranded siNA.
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ADO14515
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11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0366782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic; vasotropic; nephrotropic; cancer; restenosis; polycystic kidney disease; RNA interference; short interfering RNA; sirl short interfering nucleic acid, siNA; short interfering RNA; shRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; human; interleukin-2; ss.
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                                                                                                                                                                                                                                                                                                                                                              The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human interleukin-2 gene by RNA
                                                                                                                                                                                                                                                                                                                                                                                                                        Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            interfering nucleic acid, useful e.g. for of cancer, downregulates expression of an
                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO
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RESULT 1046
ADO14387/c
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Matches
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11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-036782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
         regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA, double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siRNA; conjugates and/or complexes of siRNA; and vectors that express siNA. The siNAs are used to modulate expression of the interleukin-2 gene in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cytostatic; vasotropic; nephrotropic; cancer; restenosis; polycystic kidney disease; RNA interference; short interfering nucleic acid; siNA; short interfering R double-stranded RNA; micro-RNA; miRNA; short hairpin RNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expression modulation; gene therapy; therapeutic target identification; pl
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                                                                                                                                                                        The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human interleukin-2 gene by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense
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                                                                                                                                                                                                                                                                                  diagnosis
                                                                                                                                                                                                                                                                                                New short
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for treating cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              function analysis; gene mapping; human; interleukin-2; ss
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                                                                                                                                                                                                                                                                                interfering nucleic acid, useful e.g. for treatment ar of cancer, downregulates expression of an interleukin
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                                                                                                                                                                                                                                                    ID NO 122; 138pp; English.
                                                                                                                                                                                                                                                                                                                                                                                          PHARM INC
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1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  y; drug screening; diagnosis; pharmacogenomics; og; human
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Example 3; SEQ ID

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260; 415pp;

English.

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RESULT 1047
AD568376/c
ID AD5683776/c
ID AD568377
XX AD56837
XX AD
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Best Local
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                                                                                                                                                                                                                      Anderson
Ellerman
Mehraban
                            New NOVX polypeptides useful for treating cancers, blood disorders, asthma, psoriasis, vascular disorders, hypertension, viral, bacterial parasitic infections, allergy, renal disorders and skin disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-FEB-2001; 2001US-0269098P.
27-FEB-2001; 2001US-0271855P.
02-MAR-2001; 2001US-0272920P.
18-APR-2001; 2001US-0284549P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-JAN-2001; 2001US-0265517P.
07-FEB-2001; 2001US-0267057P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   dermatological; osteopathic; antiarthritic; antiinflammatory; cytostatic; hypotensive; cardiant; hypertensive; antiulcer; antiallergic; antianginal; immunosuppressive; antidepressant; neurodegenerative;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Crohn's disease; rheumatoid arthritis; immunological; endocrine; pigmentation; haematopoietic; psychotic; autoimmune; muscular; osteoporosis; angina pectoris; hypotension; anxiety; allopecia; bulimia; cancer; manic depression; virucide; antibacterial; analgesic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADP68376 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human interleukin-2-targeted double-stranded siNA, which is identical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-JAN-2001; 2001US-0262454P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human; PCR; ss; NOVX; Alzheimer's disease; Huntington's; inflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-AUG-2004
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25-JAN-2001; 2001US-0264159P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer used to amplify human NOV14 DNA (Ag210) SeqID 260.
                                                                                                                                                                                                                                                                                                                                     (CURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   neuroprotective; nootropic; cerebroprotective; anticonvulsant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             interleukin-2 transcript target sequence.
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                                                                                                                                                                                    Burgess CE, Casman SJ, Colman S, Edinger S; Gerlach V, Gunther E, Kekuda R, Macdougall : Patturajan M, Rothenberg M, Shimkets RA, Smi Stone DJ, Vernet CAM, Zerhman pr
                                                                                                                                                                                                                                                                                                                                                                                      2001US-0284549P.
2001US-0285040P.
2001US-0286287P.
2001US-0303229P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 19;
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                                                                                                                                                                                                                                                     JR;
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RESULT 1048
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention relates to novel nucleic acid molecules encoding NOVX CC polypeptides selected from NOV1 to NOV11 inclusive, as well as variants CC thereof. Specifically, it refers to vectors, host cells, antibodies, CC agonists, antagonists and recombinant methods for producing proteins and composits, antagonists and recombinant methods for producing proteins according GPCRs, secretory proteins and dual specificity phosphatases. CC including GPCRs, secretory proteins and dual specificity phosphatases. CC including Crohn's disease that can be used to treat neurodegenerative cincluding Crohn's disease and rheumatoid arthritis, as well as compositions that can be used to treat neurodegenerative conditions including crohn's disease and rheumatoid arthritis, as well as conditions including osteoporosis, angina pectoris, hypotension, anxiety, CC conditions including osteoporosis, angina pectoris, hypotension, anxiety, CC anlipsecia, bulimia, cancer and manic depression. As such, they exhibit various activities including vulnerary, virucide, antibacterial, CC dermatological, osteopathic, antiarthritic, antiinflammatory, cytostatic, CC dermatological, osteopathic, antiarthritic, antiinflammatory, cytostatic, CC antianginal, immunosuppressive and antidepressant. This oligonucleotide is a PCR primer used to amplify human NOVX DNA in an exemplification of the invention.
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Best Local 9
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 This invention relates to a novel method of diagnosing a predisposition to fat deposition, leanness or non-insulin dependent diabetes mellitus (NIDDM) in a subject. The method comprises detecting the presence or
                                                                                                     Diagnosing predisposition to fat deposition, leanness or non-insulin dependent diabetes mellitus (NIDDM) comprises detecting the presence absence of a polymorphic variation in a purinergic receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human purinergic receptor P2X4-related PCR primer 59.
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                                                                     Example 3; Page 70; 154pp; English.
                                                                                                                                                                          WPI; 2004-053318/05
                                                                                                                                                                                                             Adam GIR,
                                                                                                                                                                                                                                                                                    04-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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human; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fat deposition; leanness; non-insulin
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                                                                                                                                                                                                                                                 (SEQU-) SEQUENOM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention.
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18; Conserv
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                                                                                                                                                                                                             Langdown
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                                                                                                                                                                                                                                                                                    2002US-0386012P
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Pred. No. 1.
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                                                                                                                                                                                                               Denissenko MF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               dependent diabetes mellitus;
ic; anorectic; diabetes; obes;
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Best Local &
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This invention relates to a novel method of diagnosing a predisposition to fat deposition, leanness or non-insulin dependent diabetes mellitus (NIDDM) in a subject. The method comprises detecting the presence or absence of a polymorphic variation associated with fat deposition, nucleotide sequence in a nucleic acid sample from a subject. The invention may be useful for the development of compounds with an antidiabetic or anorectic activity. The method is useful for diagnosing a predisposition to fat deposition, leanness or NIDDM. The nucleic acid encoding the polypeptide is useful for diagnosing conditions or diseases including fat deposition or NIDDM, also in treating diabetes and obesity. The present sequence is that of a PCR primer which was used for amplification or a region of the human purinargic receptor (P2X4) gene sequence in the exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Diagnosing predisposition to fat deposition, leanness or non-insulin dependent diabetes mellitus (NIDDM) comprises detecting the presence absence of a polymorphic variation in a purinergic receptor.
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human; PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        fat deposition; leanness; non-insulin dependent diabetes mellitus; NIDDM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 3; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SEQU-)
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18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               70; 154pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          receptor P2X4-related PCR primer 63.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antidiabetic; anorectic;
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Denissenko MF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Smylie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diabetes; obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ኟ
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RESULT 1050
ADH76756/c
ID ADH76756/c
XX ADH7675
XX ADH7675
XX ADH7675
XX ADH7675
XX MCHR1 g
XX melanir
XX melanir
XX WO20031
XX Unident
XX Unident
XX Unident
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                                                                                                                                                                                         The invention relates to a novel diagnostic polynucleotide composition.

The polynucleotide composition comprises: a sequence encoding a polypeptide with defined sequences given in the specification; a sequence capable of hybridizing to a melanin-concentrating hormone receptor 1

(MCHRI) gene, a polynucleotide encoding an MCHRI polypeptide; or a sequence comprising one or more of the nucleotide exchanges (SNP's) given in the specification and at least 8 bases of surrounding sequence of the CMCHRI gene. The composition has anorectic activity. The polynucleotide composition may be used in gene therapy to treat the disorders of the invention. The composition is useful for diagnosing obesity related to the presence of a molecular variant of the MCHRI gene or a susceptibility to the disorder. The MCHRI protein or polynucleotide is useful for preparing a medicament for treating or preventing obesity related to the presence of a molecular variant of the MCHRI gene. This polynucleotide is useful for presence of a molecular variant of the MCHRI gene. This polynucleotide
                                                    Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local (
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                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Platzer M,
Reichwald
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADH76756 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    obesity; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               melanin-concentrating hormone receptor 1; MCHR1; anorectic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                presence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New diagnostic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JUN-2002; 2002EP-00012569
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUN-2003; 2003WO-EP005917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-DEC-2003
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                                                                                                                                     Sequence
                                                                                                                                                                            represents an MCHR1 primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   390 AAGTGCTGGGATTACAGGC 408
             864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        UNIV PHILIPPS MARBURG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2; Page 43; 76pp;
                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of a molecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAGTGTTGGGATTACAGGC 1
GCTGGGATTACAGGCGTGA 882
                                                                                                                                     19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Platzer C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                   ₿₽;
                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    composition, useful nolecular variant of
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                                                                        1.8%;
94.7%;
                                                                                                                                     8 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        analysis primer #65.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gudermann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₽P
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Pred. No. 1.4e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G; 5 T; 0 U;
                                                  Score 17.4; D
Pred. No. 1.4e
0; Mismatches
                                                      ,
,
                                                                                                                                     G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                 the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for diagnosing obesity related to the the MCHR1 gene or a susceptibility to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hebebrand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .4e+03;
                                                                            .4e+03
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                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1;
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                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hinney
                                                                                                  19;
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                                                        Gaps
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Human interleukin-18 gene polymorphism related

probe,

SEQ ID No

58.

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RESULT 1051
ADH76751
XXXXX
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                                                                                  RESULT 1052
ADM32301/c
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                             capable of hybridizing to a melanin-concentrating hormone receptor 1 (MCHR1) gene; a polynucleotide encoding an MCHR1 polypeptide; or a sequence comprising one or more of the nucleotide exchanges (SNP's) given in the specification and at least 8 bases of surrounding sequence of the MCHR1 gene. The composition has anorectic activity. The polynucleotide composition may be used in gene therapy to treat the disorders of the invention. The composition is useful for diagnosing obesity related to the presence of a molecular variant of the MCHR1 gene or a susceptibility to the disorder. The MCHR1 protein or polynucleotide is useful for preparing a medicament for treating or preventing obesity related to the presence of a molecular variant of the MCHR1 gene. This polynucleotide represents an MCHR1 primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Platzer M,
Reichwald
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          melanin-concentrating hormone receptor 1; MCHR1; anorectic; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MCHR1 genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADH76751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADH76751 standard; DNA;
             20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel diagnostic polynucleotide composition. The polynucleotide composition comprises: a sequence encoding a polypeptide with defined sequences given in the specification; a sequence capable of hybridizing to a melanin-concentrating hormone receptor 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New diagnostic composition, useful presence of a molecular variant of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             obesity; primer;
                                                                                                                                                                                                                                                      Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Page 43; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JUN-2002; 2002EP-00012569
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JUN-2003; 2003WO-EP005917
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                                         ADM32301;
                                                                    ADM32301
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2004-062377/06
                                                                                                                                                                     731 TAGCTGGGACTACAGGCGC 749
                                                                                                                                          _
                                                                                                                                                                                                                Similarity
                                                                    standard;
                                                                                                                                                                                                                                                        BP; 5 A; 5 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Platzer C,
                                                                                                                                                                                                 Conservative
             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88
                                                                     DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                            1.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ₽₽
                                                                     ВÞ
                                                                                                                                                                                                <u>,</u>
                                                                                                                                                                                                                Pred.
                                                                                                                                                                                                                            Score 17.4;
                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hebebrand J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diagnosing
MCHR1 gene
                                                                                                                                                                                                                .4e+03;
                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                          Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          obesity related to the or a susceptibility to
                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hinney
                                                                                                                                                                                                0,
                                                                                                                                                                                                Gaps
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RESULT 1053 ADL25727/c ID ADL2572

ADL25727 standard; DNA; 19

Human NOVX gene,

forward PCR primer #29

20-MAY-2004 ADL25727;

(first entry)

Homo sapiens

autoimmune

disorder;

cancer.

ss; PCR; primer; Cytostatic; Neuroprotective; Immunosuppressive;
Gene therapy; Vaccine; human; neurodegenerative disorder;

08-JAN-2004. US2004005557-A1

16-JAN-2001; 2001US-0261376P 18-JAN-2001; 2001US-0262454P 18-JAN-2001; 2001US-0262587P

16-JAN-2002; 2002US-00051874.

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19

GCTGGTCTCCAACTCCCGA 1

210 GCTGGTCTCGAACTCCCGA 228

Conservative

0,

Indels

0

Gaps

0

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Query Match
Best Local S
Matches 18
                                                                                                                                                                                 The invention relates to a novel method for detecting a gene polymorphism in a human interleukin (II)-18 gene. The method involves detecting a 9 base insertion between -6311 position and -6310 position, a polymorphism at positions -5890, -5316, -4762, -4675, -3268, -689 and -640 of a polymucleotide which consists of a fully defined sequence of 6640 base pairs as given in the specification, where in the 6640bp polymucleotide, the position 6575 is set to +1 from which numbering is performed. The method is useful for detecting gene polymorphism in II-18 gene of human and for detecting adult onset still disease. This polymucleotide sequence represents a probe of the human interleukin-18 gene of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
Synthetic.
                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detecting gene polymorphism in interleukin-18 gene of human, useful for detecting adult onset still disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human interleukin-18; IL-18; adult onset still
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-174121/17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SUGI/) SUGIURA S.
(HYUB-) HYUBITTO GENOMICS
                                   Local Similarity
   18;
                                                                                                                          19 BP; 4 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 58; 61pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polymorphism; ss; probe
                                   94.78;
                                                                1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       즛
Score 17.4; DB 1;
Pred. No. 1.4e+03;
0; Mismatches 1;
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                                                                Length 19;
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Query Match
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14-FBB-2001; 2001US-0265595P.
28-FEB-2001; 2001US-02727409P.
16-MAR-2001; 2001US-0276777P.
17-MAY-2001; 2001US-0291672P.
27-SEP-2001; 2001US-0325306P.
18-OCT-2001; 2001US-0330336P.
09-NOV-2001; 2001US-0345202P.
                                                                                                                                                                                                     Macdougall JR, Malyankar UM, Patturajan M, Tchernev VT, Zerhusen BD, Millet I, Miller Smithson G, Baumgartner JC, Herrmann JL, P Mezes PD, Kekuda R, Taupier RJ, Gerlach V, Ellerman K, Rothenberg M, Stone DJ, Burges
                                                        The invention relates to novel human NOVX nucleic acids and polypeptides. The polypeptide, nucleic acid or antibody is useful for preparing a composition for treating or preventing a NOVX-associated disorder, e.g., neurodegenerative or autoimmune disorders or cancer. The present sequence represents a PCR primer used to isolate human NOVX genes of the
                                                                                                                                         New NOVX polypeptide, useful for preparing a preventing a NOVX-associated disorder, e.g., autoimmune disorders or cancer.
                                                                                                                                                                                                                                                          Padigaru M,
Vernet CAM,
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(GORM/)
(MEZE/)
(KEKU/)
(TAUP/)
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(BAUM/)
(HERR/)
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(PATT/)
(SHIM/)
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(PENA/)
(TCHE/)
(ZERH/)
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(SHEN/)
(CASM/)
(GUOX/)
(EDIN/)
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(ALSO/)
(COLM/)
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(GROS/)
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                             Sequence
                                                                                                                      Example
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//) BOLDOG F L.
//) VERNET C A M.
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MALYANKAR U M.
PATTURAJAN M.
SHIMKETS R A.
PENA C E A.
TCHERREV V T T.
ZERHUSEN B D.
MILLET I.
MILLET G E.
LEPLEY D M.
SMITHSON G.
BAUMGARTNER J C.
                                                                                                                                                                                                                                                                                                                                           TAUPIER R J.
GERLACH V.
GROSSE W M.
                                                                                                                                                                                                                                                                                       ELLERMAN K.
ROTHENBERG M.
STONE D J.
BURGESS C E.
                                                                                                                                                                                                                                                                                                                                                                                                     HERRMANN J 1
PEYMAN J A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SHENOY S G.
CASMAN S J.
                                                                                                                     3; Page 263; 282pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                GORMAN L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EDINGER S
                                                                                                                                                                                                                                                                                                                                                                           KEKUDA R.
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                              B₽;
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Li L, Shenoy
                                                                                                                                                                                                                       Isobrook JP, Colman SD, Spytek KA, Boldog FL; i L, Shenoy SG, Casman SJ, Guo X, Edinger SR; Malyankar UM, Patturajan M, Shimkets RA, Pen Zerhusen BD, Millet I, Miller CE, Lepley DM; Cerhusen BD, Millet I, Peyman JA, Gorman JA, Tin
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1.8%;
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Score 17.4; DB 1;
Pred. No. 1.4e+03;
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lach V, Grosse WM,
Burgess CE;
                              0 Other;
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         Length
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Best Local Similarity

ADP09402
ID ADPO
XX
AC ADPO
XX
DT 26-1

standard; DNA;

19

ВP

26-AUG-2004 ADP09402; ADP09402

(first entry)

RESULT 1055

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ADPO8706/c
ID ADPO87
XX ADPO87
XX ADPO87
XX ADPO87
XX Extend
XX breast
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying a subject at risk of breast cancer by detecting the presence or absence of one or more nucleotide polymorphic variations, useful for diagnosing, preventing and/or treating breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-NOV-2002; 2002US-0429136P.
24-JUL-2003; 2003US-0490234P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GP6; GPIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Extend primer 43 used to genotype human glycoprotein VI polymorphism.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    breast cancer;
                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                             nucleotide
GPIV;GPVI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-441082/41.
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                                              377 CCTCAGCCTCCCAAAGTGC
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                                                                                                                      Similarity
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CCTCAACCTCCCAAAGTGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 82;
                                                                                                                                                                                                                                             polymorphisms within human glycoprotein VI (platelet) DNA which is located at chromosomal position 19q13.4.
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                                                                                                                                                                                               3 A;
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                                                                                                                    1.8%;
94.7%;
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                                                                                                                      Score 17.4; DB 1;
Pred. No. 1.4e+03;
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RESULT 1056
AD080022
ID AD08002
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AC AD08002
XX CENPC1
XX CENPC1
XX CENPC1
XX CENPC1;
KW CENTON
XX CENTON
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XX PD 10-JUN-
XX 25-NOV-
PF 25-NOV-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel method for identifying a subject at risk of breast cancer which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a risk of breast cancer, as well as therapeutic and prophylactic treatments that specifically target breast cancer, such as gene therapy. The current sequence is that of a Extend primer of the invention which was used to genotype single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                or absence of diagnosing,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 3 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identifying a subject at risk of breast cancer by detecting the presence or absence of one or more nucleotide polymorphic variations, useful for diagnosing, preventing and/or treating breast cancer.
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                 25-NOV-2003; 2003WO-US037943.
                                                                                                                                   Cytostatic; Gene therapy; breast cancer; human; DLG1; KIAA0783; CENPC1; SNP; single nucleotide polymorphism; centromere protein Centromere autoantigen C1; chromosome 4q12-q13.3; extend; primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 6; Page 110; 286pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-NOV-2002; 2002US-0429136P
24-JUL-2003; 2003US-0490234P
                                                                          WO2004047514-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleotide
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                                                10-JUN-2004
                                                                                                                                                                                                                                                            AD080022
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                                                                                                                                                                                                                                                                                                                                                                CTCTGTCACCCATGCTGGA 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polymorphisms within human LOC338749 DNA which is position 11p15.3.
                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                         DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                          94.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            to genotype human LOC338749 polymorphism
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene therapy; human; LOC338749;
R; primer; SNP; single nucleotide polymorphism
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 17.4;
Pred. No. 1.
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The present invention relates to a method for identifying a subject at CC risk of breast cancer. The method comprising detecting the presence or cc absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The nucleic acid sample (comprises the DLG1 region (AD079402), KIAA0783 region (AD079403), DPF3 (cromprises the DLG1 region (AD079402)). The gene DLG1 (discs, region (AD079404) or CENPC1 region (AD079405). The gene DLG1 (discs, pr. hdlg or SAP97. DLG1 has been mapped to chromosomal position 3q29. The gene KIAA0783 is also known as synapse-associated protein (cromprises as transcription factor. The gene protein 14. KIAA0783 and protein is a convel gene with unknown function, however, being a zinc finger protein, crit likely to be a transcription factor. The gene DPF3 (D4, zinc and couble PHD fingers, family 3) is also known as CERD4, cer-d4, FLJ14079 and 2810403B03Rik. DPF3 is a Rho family guanine-nucleotide exchange factor. DPF3 has been mapped to chromosomal position 14q24.3-q31.1. The CENPC1 has been mapped to chromosomal position 4q12-q13.3. CENPC1 is a centromere autoantigen and a component of the inner chinecochore size and a timely transition to anaphase. The method is useful for identifying a subject at risk of breast cancer, to analyze and component sequence was used in an example from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying a subject at risk of breast cancer by detecting the presence of polymorphic variations in the DLG1, KIAA0783, DPF3 or CENPC1 regions which are associated with breast cancer in a nucleic acid sample from a
Sequence 19 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-NOV-2002; 2002US-0429136P.
24-JUL-2003; 2003US-0490234P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 6; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 subject
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-441037/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Roth RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENOM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nelson MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     91; 227pp; English.
A; 7 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Braun
  G; 6 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kammerer SM, Reneland
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              S
                                Query Match
Best Local Similarity
Matches 18; Conserv
                970
\vdash
           TCGGCTCACTGCAACCTCT 988
                                 Conservative
                                         1.8%;
                                 <u>,</u>
                                Score 17.4; D
Pred. No. 1.4e
0; Mismatches
                                          1.4e+03
                                                   DB 1;
                                  1;
                                                  Length 19;
                                  0
                                  Gaps
                                  0
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RESULT 1057
AAZ07267/c
ID AAZ0726
Homo
                                                   Synthetic.
                                                                                                     Human telomerase RNA
                                                                                                                       22-OCT-1999
                                                                                                                                        AAZ07267;
                                                                                                                                                         AAZ07267 standard; DNA; 20
                                                                          Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human;
                                                                  primer;
                                         sapiens
                                                                    88
                                                                                                                      (first entry
                                                                                                     gene
                                                                                                     (hTR)
                                                                                                     specific primer hTR10F
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05-AUG-1999

WO9938964-A2

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RESULT 1058
AAZ37719/c
ID AAZ3771
XX
AC AAZ3771
XX
DT 07-JAN-
XX
DE Human 1
XX
Human 1
XX
Human 1
XX
Human 2
XX
Human 5
XX
Human 6
XX
Human 6
XX
Human 7
XX
Human 7
XX
Human 7
XX
Human 1
X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC heterologous gene, especially a gene encoding a cytotoxin, for therapy of crancer, especially neoplasias. The telomerase is necessary for the CC unrestricted proliferative capacity of many human cancers. Mutation or CC dysregulation of the telomerase repression pathway may cause reactivation CC or upregulation of the telomerase expression in cancer. Substances, CC identified in the methods, can be used to block transcription from the TR CC gene promoter through interaction of the 5' regulatory sequences. These CC gene promoter therapy. In particular, gene therapy vectors CC useful for cancer therapy. In particular, gene therapy vectors CC (especially pGT62-codAupp) comprising the promoter and a viral thymidine CC (inase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of CC telomerase RNA gene through manipulation of transcription factors may be ceffective anticancer therapy and the cloning of the hTR gene promoter CC allows the analysis of therapeutic molecules which modulate hTR promoter CC telomerase therapy and the cloning of the hTR gene promoter CC activity. Sequences AAZO7623-80 represents PCR primers for amplifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse and human telomerase RNA gene gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human mdm2 phosphorothioate oligodeoxynucleotide #249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ37719 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Fig 6; 109pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CANC-) CANCER
                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                   Human mdm2 gene; proliferation; tumour; phosphorothioate; antisense; modulation; oligonuclectide; expression; inhibi antisense; brodulation; blood cancer; brain cancer; breast can hyperproliferation; blood cancer; psoriasis; fibrosis; athe lung cancer; soft tissue cancer; psoriasis; fibrosis; athe
                26-MAR-1998;
                                                                           26-MAR-1999;
                                                                                                                                         30-SEP-1999
                                                                                                                                                                                                      WO9949065-A1
                                                                                                                                                                                                                                                                                                                                                         restenosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention relates to promoter regions from mouse and human telomerase (TR) component genes. The TR gene promoter can be linked to a
                                                                                                                                                                                                                                                              sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              717
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CTCAGCCTCCTGAGTAGCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CCCAGCCTCCTGAGTAGCT 735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene (hTR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP;
                                                                                                                                                                                                                                                                                                                                                             88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98GB-00001902
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99WO-GB000308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6 A;
                                                                           99WO-US006702
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   promoter sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.8%;
94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>.</u>.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 17.4; DB 1
Pred. No. 1.5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             promoters, useful for tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                           inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     p53; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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RESULT 1059
AAZ37727/c
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Query Match
Best Local Sim
Matches 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treat an animal having a disease or condition associated with mdm2, particularly a hyperproliferative condition, more particularly cancer, especially of the blood, brain, breast, lung or soft tissue, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         novel nucleotide antisense compounds, targetted to the 5' untranslated translation termination codon, or 3' untranslated region of a nucleic acid encoding human mdm2, that modulates expression of human mdm2. The oligonucleotides mediate their effect by antisense inhibition of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ37473-Z37738 represent human mdm2 phosphorothioate oligonucleotides. AAZ37471, AAZ37472, AAZ37739, AAZ37740 and AAZ37741 are used in the exemplification of the present invention. The present invention describes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Miraglia LJ,
                                                                                                                                                                                                                                                                                                                Human mdm2 gene; proliferation; tumour; phosphorothicate; p53; cancer; antisense; modulation; oligomucleotide; expression; inhibition; hyperproliferation; blood cancer; brain cancer; breast cancer; ancer; sencer; sencer; sencer; sencer; sencer; sencer; procession; lung cancer; soft tissue cancer; psoriasis; fibrosis; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 6 A; 2 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compounds used to treat
                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                Human mdm2 phosphorothioate oligodeoxynucleotide #257
                                                                                                                                                                                                                                                                                                                                                                                                           07-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ37727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hyperproliferative gene expression. The antisense compound
                                                                            WPI; 1999-610754/52
                                                                                                   Miraglia LJ,
                                                                                                                                                       26-MAR-1998;
                                                                                                                                                                                                          30-SEP-1999.
                                                                                                                                                                                                                                   WO9949065-A1
                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS
                                                                                                                                                                                26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    531 CATCCTCCTGCCTCAGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9; Page
                                                                                                                              SISI
                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CATTCTCCTGCCTCAGCCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fibrosis, atherosclerosis or restenosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PHARM
                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nero P,
                                                                                                                              PHARM INC
                                                                                                   Nero P,
                                                                                                                                                        98US-00048810
                                                                                                                                                                                 99WO-US006702
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  54; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Graham MJ,
                                                                                                      Graham MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17.4;
No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Monia BP,
                                                                                                       Monia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            eg. hyperproliferative conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .5e+03;
                                                                                                       B₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B
                                                 hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cowsert
                                                                                                       Cowsert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       untranslated,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    is used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                     conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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AAZ37473-Z37738 represent human mdm2 phosphorothioate oligonucleotides.

Example

9; Page

55;

157pp; English

New antisense compounds used

to treat

eg.

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RESULT 1060
RAZ37726/c
ID AAZ3772
XX AAZ3772
XX AAZ3772
XX Human m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              exemplification of the present invention. The present invention describe novel nucleotide antisense compounds, targetted to the 5' untranslated, translation termination codon, or 3' untranslated region of a nucleic acid encoding human mdm2, that modulates expression of human mdm2. The oligonucleotides mediate their effect by antisense inhibition of hyperproliferative gene expression. The antisense compound is used to treat an animal having a disease or condition associated with mdm2, particularly a hyperproliferative condition, more particularly cancer, especially of the blood, brain, breast, lung or soft tissue, or psoriasis, fibrosis, atherosclerosis or restenosis
                            hyperproliferative gene expression. The antisense compound is used to treat an animal having a disease or condition associated with mdm2, particularly a hyperproliferative condition, more particularly cancer, especially of the blood, brain, breast, lung or soft tissue, or psoriasis, fibrosis, atherosclerosis or restenosis
                                                                                                                                                                                           AAZ37473-Z37738 represent human mdm2 phosphorothioate oligonucleotides.
AAZ37471, AAZ37472, AAZ37739, AAZ37740 and AAZ37741 are used in the exemplification of the present invention. The present invention describes novel nucleotide antisense compounds, targetted to the 5' untranslated, translation termination codon, or 3' untranslated region of a nucleic acid encoding human mdm2, that modulates expression of human mdm2. The oligonucleotides mediate their effect by antisense inhibition of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human mdm2 gene; proliferation; tumour; phosphorothicate; p53; cancer; antisense; modulation; oligonuclectide; expression; inhibition; hyperproliferation; blood cancer; brain cancer; breast cancer; antisers cancer; section; brosses; atherosclerosis; lung cancer; soft tissue cancer; psoriasis; fibrosis; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 9 A; 4 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Miraglia LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9949065-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JAN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ37726 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compounds used to treat eg. hyperproliferative conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human mdm2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1999-610754/52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9; Page 55; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TTTGTATTTTTAGTAGAGA 789
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ37472,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     phosphorothioate oligodeoxynucleotide #256.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nero P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98US-00048810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99WO-US006702
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.8%;
94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ37739, AAZ37740 and AAZ37741 are used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Graham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention describes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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RESULT 1061
AAZZ1805/c
ID AAZZ180
XX AAZZ180
XX O1-DEC-
XX Neempla
XX neeppla
XX neeppla
XX Neempla
XX OUYJO)
XX Sidran
XX OUYJO)
XX Sidran
XX OUYJO)
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XX Detect
CC Neempla
CC Oneopla
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CC Ceteloc
CC Gevelor
CC Geneopla
CC This a
CC Chemop
XX Sequen
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                                                           Query Match
Best Local Similarity
Matches 18; Conserv
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                                                                                                                                                                  This is an exemplary oligonucleotide primer, for use in the detection of neoplasmic related gene mutations. There are over 40 known proto-
oncogenes and suppressor genes to date, which control growth,
development, and cell differentiation. Regulation of these genes can,
under certain circumstances, be altered and normal cells can assume
neoplastic growth characteristics. The invention provides a method for
detecting a neoplastic disorder of the head and neck or lung in a
subject. The detection of a target mutant nucleotide sequence in the
saliva is indicative of a neoplastic disorder of the head, neck or lung.
This allows early detection and therefore treatment of the preneoplasia
or cancer, and can also be used to monitor high risk patients undergoing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       neoplasia; mutant; target nucleotide; hybridization; lung cancer; ss; neck cancer; head cancer; saliva test; chemotherapy; early detection; primer; PCR; amplification.
                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Exemplary oligonucleotide primer X80250 (For).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ21805;
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                                                                                                                                                        chemoprevention
                                                                                                                                                                                                                                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                   Detection of cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-551428/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9946408-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYJO ) UNIV JOHNS HOPKINS SCHOOL MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
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                             646
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 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 BP; 4 A; 3 C; 8
 AGGCTGGAGTGCAGTGGTG 2
                             AGGCTGGAGTGCAGTGGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CCACCACACCTGGCTAATT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCACTACACCTGGCTAATT 596
                                                                                                                            20 BP; 4
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                                                                                                                                                                                                                                                                                                                                                                    Page 29;
                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-00038637
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                                                                                                                                                        or chemotherapy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.8%;
94.7%;
                                                                            1.8%;
                                                                                                                            10 C;
                                                                                                                                                                                                                                                                                                                                                                   99pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                     comprises assaying
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                                                                                                                            2 G; 4 T;
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Pred. No. 1
                               664
                                                                              Pred.
                                                                                           Score 17.4; DB 1; Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5
H;
                                                                Mismatches
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                                                                              No.
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                                                                                                                            0
U;
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                                                                                                                                                                                                                                                                                                                                                                                                                     for a genetic mutation associated
                                                                            .5e+03
                                                                                                                            0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 20;
                                                                Indels
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                                                               Gaps
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RESULT 1062 AAF31821/c

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RESULT 1063
AAF80881/c
ID AAF8088
XX AAF8088
XC AAF8088
XC AAF8088
XX O2-MAY-
XX Human T
XX Antiser
XX Antiser
XX Antiser
XX US61842
XX US61842
XX O6-FEB-
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                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                     nucleobases in length that have been designed to target a 5'untranslated region, start codon, coding region or 3'untranslated region of the human receptor activator of NF kappaB (RANK). The antisense compounds specifically hybridise with and inhibit the expression of RANK. The antisense oligonucleotides are useful for inhibiting the expression of human RANK in human cells or tissues. They can be utilised for diagnostics, therapeutics for the treatment of diseases associated with the expression of RANK, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, and as research reagent. The antisense compounds are safely and effectively administered to humans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human RANK antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF31821 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel antisense compounds capable of modulating expression of human receptor activator of NF-kappaB useful for diagnosis, prophylaxis attreatment of diseases associated with expression of RANK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-APR-2001
                                                                                                                                                                                    AAF80881 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 4 A; 3 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 14; Col 44; 40pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    receptor activator
                                                                                                                                 02-MAY-2001
                                                                                                                                                          AAF80881;
     06-FEB-2001
                           US6184212-B1
                                                     Homo sapiens
                                                                               Antisense; mdm2;
                                                                                                                                                                                                                                                                            1111
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                                                                                                                                                                                                                                                  19
                                                                                                         mdm2
                                                                                                                                                                                                                                                                                                      18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cytostatic;
                                                                                                                                                                                                                                                                            CAGGCTGGTCTCAAACTCC 1129
                                                                                                                                                                                                                                                  CAGCCTGGTCTCAAACTCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cowsert LM;
                                                                                                       phosphorothicate oligonucleotide
                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-00435296
                                                                               hyperproliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antiinflammatory; antisense oligonucleotide; cancer;
of NF-kappaB; RANK; infection; inflammation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                 entry)
                                                                                                                                                                                                                                                                                                                  1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English.
                                                                                                                                                                                    20
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                                                                                                                                                                                    ВР
                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                    Score 17.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                cancer; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO:
                                                                                                                                                                                                                                                                                                                     1.5e+03
                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                          #255
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                                                                                                                                                                                                                                                                                                                                Length 20;
                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                       Gaps
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RESULT 1064
AAF80873/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to an antisense compound 8-30 nucleobases in length targeted to nucleobases 1-308 of the 5' untranslated region, 1776-1806 of the translation termination codon region or 1818-2370 of the 3' untranslated region of a nucleic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro. The hyperproliferative disorder includes cancer or psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel antisense compound 8-30 nucleobases in length targeted to a nucleoacid molecule encoding human mdm-2 useful for modulating the expression of human mdm-2 and reducing hyperproliferation of human cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 9 A; 4 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Miraglia LJ,
                                                                 Novel antisense compound 8-30 nucleobases in length targeted to a nucleic acid molecule encoding human mdm-2 useful for modulating the expression of human mdm-2 and reducing hyperproliferation of human cells.
                                                                                                                                                                                                                                                                                                                                Human mdm2
                                                                                                                                                                                                                                                                                                                                                        02-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                              AAF80873;
                                                                                                                                                                                                                                                                                                                                                                                                     AAF80873
                                                                                                                                       Miraglia LJ,
                                                                                                                                                                                      26-MAR-1998;
                                                                                                                                                                                                             26-MAR-1999;
                                                                                                                                                                                                                                     06-FEB-2001.
                                                                                                                                                                                                                                                           US6184212-B1
                                                                                                                                                                                                                                                                                   Homo
                                                                                                                                                                                                                                                                                                         Antisense; mdm2;
                                                                                                                                                              (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001-190948/19
                                                                                                                                                                                                                                                                                  sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            771 TTTGTATTTTTAGTAGAGA 789
                                                                                                                                                                                                                                                                                                                                                                                                                                                              20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                TTTGTACTTTAGTAGAGA
                                                                                                                                                                                                                                                                                                                                phosphorothicate oligonuclectide #247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PHARM INC
                                                                                                                                                                PHARM INC.
                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                        Nero P,
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                                                                                                                                                                                                             99US-00280805
                                                                                                                                                                                                                                                                                                           hyperproliferation; cancer; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94.7%;
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                                                                                                                                         Monia
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                                                                                                                                         B₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
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The present invention relates to an antisense of in length targeted to nucleobases 1-308 of the 1776-1806 of the translation termination codon

compound 8-30 nucleobases e 5' untranslated region, n region or 1818-2370 of the

lated region, 1818-2370 of the

Example

9; Col

31; 77pp;

English.

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RESULT 1065
AAF80880/c
ID AAF8088
AAH40109
ID AAH
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Best Local (
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                                                                                                                                                                                                                                                                                                                        Novel antisense compound 8-30 nucleobases in length targeted to a nucleic acid molecule encoding human mdm-2 useful for modulating the expression of human mdm-2 and reducing hyperproliferation of human cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3' untranslated region of a nucleic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro.
                                                                                                                                                                                                           The present invention relates to an antisense compound 8-30 nucleobases in length targeted to nucleobases 1-308 of the 5' untranslated region, 1776-1806 of the translation termination codon recording human mdm-2.370 of the 3' untranslated region of a nucleic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human mdm2
                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                    Miraglia LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense; mdm2; hyperproliferation; cancer; psoriasis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF80880;
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                                                                                                                                                                                                                                                                                                   Example
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                                                                                                                                                                                                   hyperproliferative disorder includes cancer or
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                                                                                                 578 CCACTACACCTGGCTAATT 596
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                                                                                                                                       Similarity
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                                                                                                                                                                          BP; 4 A; 3 C; 8 G; 5 T; 0 U; 0 Other;
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                                                                                                                          Conservative
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                                                                                                                                    1.8%;
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Pred.
                                                                                                                                       Score 17.4;
Pred. No. 1.
                                                                                                                                      Pred.
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                                                                                                                          Mismatches
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No. 1.
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                                                                                                                                      1.5e+03;
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                                                                                                                                                   DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                     Cowsert LM;
                                                                                                                                                Length 20;
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AAH40109 standard;

DNA;

20

ВP

RESULT 1067 AAC86127 ID AAC8612

AAC86127 standard; cDNA; 20

ВP

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                                                                        Query Match
Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                        SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch.Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Single nucleotide polymorphism; SNP; single nucleotide primer SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; car
                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequences AAH37205 - AAH40944 represent PCR primers,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 64; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acid sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-290930/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-OCT-2000; 2000WO-US028436.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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                                     ATCCTCCTGCCTCAGCCTC
                                                                                                                                                BP;
                                                                                                                                                                                          SNP
                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               upper PCR primer SEQ ID
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                                                                                                                                                  3 A;
                                                                                                                                                                                        containing DNA sequence
                                                                                            1.8%;
                                                                                                                                                  9 C; 2
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                                                                                                                                                    G; 6 T; 0 U; 0 Other;
                                   550
                                                                                            Score 17.4;
Pred. No. 1.
 20
                                                                            Mismatches
                                                                                              .5e+03
                                                                                                               DB 1;
                                                                                                           Length 20;
                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single nucleotide
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                                                                            Gaps
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                                                                                                                                                                                                          glycosylation site, a transmembrane domain, and a cytoplasmic domain CQ glycosylation site, a transmembrane domain, and a cytoplasmic domain having at least one SH2-binding motif. APEX proteins and antibodies are useful in the study, diagnosis, prevention and treatment of disease associated with the presence of an APEX protein e.g., asthma, arteriosclerosis, AIDS, cirrhosis, Crohn's disease, atopic dermatitis, called the continuous anaemia, burstitis, inflammatory bowel disease, multiple cemphysema, atrophic gastritis, inflammatory bowel disease, multiple sclerosis, mysathenia gravis, myocardial or pericardial inflammation, csclerosis, mysathenia gravis, myocardial or pericardial inflammation, cancer, immune disorders, autoimmune diseases, graft rejections, graft versus host reaction and systemic lupus crythematosus. APEX proteins are useful as diagnostic and/or prognostic markers on APCs or APEX expressing cells, the ability to elicit the generation of antibodies and as targets for various therapeutic
                                                                      Matches
                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inflammatory bowel disease; multiple sclerosis; myasthenia gravis; myocardial inflammation; pericardial inflammation; osteoarthritis; osteoporosis; psoriasis; Reiter's syndrome; rheumatoid arthritis; inflammation; cancer; autoimmune disease; graft rejection; amplify; graft versus host disease; systemic lupus erythematosus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         extracellular domain; immunoglobulin-like domain; Ig-like struci
N-glycosylation site; transmembrane domain; cytoplasmic domain;
SH2-binding motif; asthma; arteriosclerosis; AIDS; cirrhosis; SH2-binding motif; asthma; arteriosclerosis; AIDS; cirrhosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Primer JNF15 to isolate APEX cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAC86127;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The sequences given in AAC86117-42 are primers which were used to isolate the cDNA sequences which encode antigen presenting cell expression (APEX)-1, APEX-2 and APEX-3 proteins. APEX-1 and APEX-2 comprise an extracellular domain having one immunoglobulin (Ig)-like structure and N-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Crohn's disease; atopic dermatitis; autoimmune anaemia; bursitis; cholecystitis; diabetes mellitus; emphysema; atrophic gastritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 50; Page 83; 112pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disease and atopic dermatitis.
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                                                                                                                                             Sequence
                                                                                                                                                                                                modalities.
                                    967
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bind APEX
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                                                                                                                                             Α;
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                                                                                     1.8%;
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immunoglobulin-like domain; Ig-like structure
                                                                                                                                               N
                                                                                                                                                                                                are also useful for identifying
                                                                      0;
                                                                                                                                             G; 4 T; 0 U; 0 Other;
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 20
                                                                                       Score 17.4;
Pred. No. 1
                                                                          Mismatches
                                                                                       .5e+03
                                                                                                            DB 1;
                                                                                                        Length 20;
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SLC6A4;

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RESULT 1068
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ID AAS0123
XX AAS0123
XX PASO123
XX Human E
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XX CURA-)
XX Homo Ba
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20-SEP-1999;
13-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The sequence represents the Reverse PCR primer, used in expression analysis of human secreted protein, POLYS, POLYX nucleic acids, polypeptides and antibodies to POLYX can be used for treating or preventing a POLYX associated disorder in a subject, preferably a human. These can be used in the manufacture of a medicament for treating a syndrome associated with a human disease selected from a POLYX-associated disorder, where the therapeutic is a POLYX polypeptide, a POLYX nucleotide or a POLYX antibody. They may also be used to screen for a modulator of activity, or latency, or predisposition to a POLYX-associated disorder account of activity, or latency, or predisposition to a POLYX-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAS01235 standard; cDNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reverse PCR primer, used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAS01235;
Solute carrier family 6 neurotransmiter transporter; genotyping; allele specific oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New POLYX polypeptide useful for associated disorder, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-SEP-2000; 2000WO-US025106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-MAR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200119856-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted
                                                                                                                             30-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 associated disorder, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-244781/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-SEP-2000;
                                                                               Primer
                                                                                                                                                                              AAF74118
                                                                                                                                                                                                                          AAF74118 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                 1000 TCAAGCGATTCTCCTGTCT 1018
                                                                                                                                                                                                                                                                                                                                                  19
                                                                                                                                                                                                                                                                                                                                                                                                                                                    18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5; Page 111; 152pp;
                                                                                                                                                                                                                                                                                                                                                     TCAAGCGATTCTCCTGCCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 7 A; 3 C; 7 G; 3 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000US-00659634
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fernandes
                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein; therapeutic; diagnostic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0154762P
99US-0159231P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0153629P.
99US-0154520P.
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                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer
                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             expression analysis of POLYS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Herrmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 17.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. 1.5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    treating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       preventing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yang
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                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to a polymorphic variant of a reference sequence for the solute carrier family 6 neurotransmitter transporter, serottonin member 4 (SLC6A4) gene or a fragment of it or a sequence complementary to the first sequence. The invention is used in producing a recombinant organism that can be used to express SLC6A4 for protein structure analysis and binding studies. A composition comprising a genotyping oligonucleotide is used to detect a polymorphism in the SLC6A4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated polynucleotide comprising a polymorphic variant for the solute carrier family 6 neurotransmitter transporter, serotonin member 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo
                               modified_base
                                                                                                                                                                                                                                                                   AAH20696;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Denton
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200109161-A1
                                                                                                                                                                                                                       Human telomeric repeat binding factor 2 oligonucleotide 111424
                                                                                                                                                                                                                                                                                         AAH20696 standard;
                                                                         modified_base
                                                                                                                    modified_base
                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                        Antisense; inhibitor;
                                                                                                                                                                                                                                               13-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 36; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GENA-), GENAISSANCE PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sapiens
                                                                                                                                                                                                                                                                                                                                                                    373 CCTGCCTCAGCCTCCCAAA 391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for identifying drugs for treating disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RR,
                                                                                                                                                                                                                                                                                                                                               N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein.
                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                      phosphorothioate; human; telomeric repeat binding factor 2;
premature aging; hyperproliferative disorder; cancer;
                                                                                                                                                                              88
                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Duda A,
                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99US-0146290P.
                                 /note= '
                                                                                                                                 Location/Qualifiers
                                                                /*tag=
                                                                                      note=
                       *tag=
                                                        mod.
                                                                                                 mod_base= OTHER
                                                                                                             *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                     A;
                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                  1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nandabalan
            base= OTHER
                                          _base= OTHER
== "2-O-methoxyethyl"
                                                                                                                                                                                                                                                                                                                                                                                                                                     9 C; 3
                                                                                      "phosphorothioate backbone"
"2-0-methoxyethyl"
                        a
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                                                                                                             σ
                                                                                                                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                   G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                               20
                                                                                                                                                                                                                                                                                                                                                                                                   Score 17.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                    .5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Stephens
                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         related to expression
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AAS29495/
ID AAS2
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           in length targeted to a polynucleotide encoding human telomeric repeat binding factor 2 (II) which specifically hybridizes with, and inhibits the expression of (II). (I) is useful for treating a human having a disease or condition associated with (II) such as premature aging or a hyperproliferative disorder especially cancer, by inhibiting the expression of (II) in human cells or tissues. (I) is useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. The products of the invention have cyrostatic activity. This sequence represents an antisense oligonucleotide used to illustrate the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense compounds targeted to nucleic acid encoding telomeric repeat binding factor 2 useful for treating conditions such as premature agin and diseases such as cancer.
                                                                                                                                                                                             Homo
                                                                                                                                                                                                                        Human; mdm2; hyperproliferative disorder; cancer; psoriasis; atherosclerosis; tumour; cytostatic; anti psoriatic; anti arteriosclerotic; vasotropic; antisense; phosphorothioa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                               Human mdm2 antisense oligonucleotide 31470.
                                                                                                                                                                                                                                                                                                             21-NOV-2001
                                                                                                                                                                                                                                                                                                                                         AAS29495;
                                                                                                                                                                                                                                                                                                                                                                     AAS29495 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel antisense compound (I) 8-30 nucleobases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Page 81; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-398071/42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200143752-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC.
                                                    US2001016575-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention
                                                                                                                                                                                            sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   969
                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 Similarity
18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTCGGCTCACTGCAACCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                        CTCGGCTCACTGCGACCTC 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000WO-US033954
                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                               /note= "OTHER= All phosphorothioate linkages,
additionally bases 1-6 and bases 15-20 are 2'-0-
methoxyethyl bases, and bases 7-14 are deoxynucleotides"
                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.8%;
                                                                                                                         base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               °.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 17.4; DB 1; Length Pred. No. 1.5e+03; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    987
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 Other;
                                                                                                                                                                                                                         phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                premature aging
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02-JAN-2001; 2001US-00752983

23-AUG-2001

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RESULT 1072
AAS29488/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             length targeted to the 5' untranslated region, coding region or translation termination codon region, 3' untranslated region, coding region or translation start site of a nucleic acid encoding human mdm2, where the antisense compound modulates the expression of human mdm2. The antisense oligonucleotides of the invention are useful for encoding human mdm2 and for inhibiting the expression of human mdm2. They may be used for treating an animal having a disease or condition associated with amplification of mdm2 gene or overexpression of mdm2 e.g. a hyperproliferative disorder such as cancer (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis, fibrosis, atherosclerosis or restenosis, tumours, colorectal carcinoma and chronic myelogenous leukemia. The antisense compound may be administered with a chemotherapeutic agent to overcome drug resistance. The antisense compound may be antisense compound reduces hyperproliferation of human cells. The method, which involves the use of the antisense compound, is also useful collections and physiological processes and useful in both clinical research and chiagnostic tools. AAS29242-AAS2907 represent the human mdm2 antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                         Best
                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           An antisense compound, useful for treating e.g. nucleobases targeted a region (e.g. translation of a nucleic acid encoding human mdm2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-MAR-1998;
26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-535565/59.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Miraglia LJ, Nero P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (NERO/)
(GRAH/)
                                                                                                                                  Human; mdm2; hyperproliferative disorder; cancer; psoriasis; atherosclerosis; tumour; cytostatic; anti psoriatic; anti arteriosclerotic; vasotropic; antisense; phosphorothioate; ss.
                                                                                                                                                                                             Human
                                                                                                                                                                                                                                                                                     AAS29488 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     oligonucleotides of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to antisense compounds, 8-30 nucleobases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MIRA/)
                                                      modified_base
                                                                                                                                                                                                                           21-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                     sapiens
                                                                                                                                                                                                                                                                                                                                                                                              578
                                                                                                                                                                                                                                                                                                                                                                  19
                                                                                                                                                                                              mdm2
                                                                                                                                                                                                                                                                                                                                                                                                                            18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MIRAGLIA L J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MONIA B P.
COWSERT L M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GRAHAM M J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                                                                             CCACTACACCTGGCTAATT 596
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                                                                                                                                                                                             antisense oligonucleotide 31623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 4 A;
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                                                                                                                                                                                                                           (first entry)
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99US-00280805.
/mod_base= OTHER
/note= "OTHER= All phosphorothioate linkages,
additionally bases 1-6 and bases 15-20 are 2'-O-
                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                           1.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 17.4;
Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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                                                                                                                                                                                                                                                                 RESULT 1073
AAS29496/c
                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-535565/59
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GRAH/)
(MONI/)
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26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-JAN-2001; 2001US-00752983
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                                    Human; mdm2; hyperproliferative disorder; cancer; psoriasis; atherosclerosis; tumour; cytostatic; anti psoriatic; anti arteriosclerotic; vasotropic; antisense; phosphorothios
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 9; Page 18; 81pp; English.
                                                                                                                      Human mdm2 antisense oligonucleotide 31627.
                                                                                                                                                                21-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to antisense compounds, 8-30 nucleobases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (NERO/)
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Homo sapiens
                                                                                                                                                                                                                                                AAS29496 standard; DNA; 20
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GRAHAM M J.
MONIA B P.
COWSERT L M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                     CATTCTCCTGCCTCAGCCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 6 A; 2 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                (first entry)
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99US-00280805.
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(e.g. translation terminat
                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           .5e+03;
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                                           phosphorothioate; ss.
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RESULT 1074
ABS67842/c
ID ABS6784
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AC ABS6784
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DT 29-NOV-
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                                                                                                                                                     문
                                                                                                                                                                                                                                                                                                                                   The present invention relates to antisense compounds, 8-30 nucleobases in clength targeted to the 5' untranslated region, translation termination codon region, 3' untranslated region, coding region or translation start site of a nucleic acid encoding human mdm2, where the antisense compound modulates the expression of human mdm2. The antisense oligonucleotides of the invention are useful for encoding human mdm2 and for inhibiting the expression of human mdm2. They may be used for treating an animal having a disease or condition associated with amplification of mdm2 gene or overexpression of mdm2 e.g. a hyperproliferative disorder such as cancer (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis, fibrosis, atherosclerosis or restenosis, tumours, colorectal carcinoma and chronic myelogenous leukemia. The antisense compound may be administered with a chemotherapeutic agent to overcome drug resistance. The antisense compound reduces hyperproliferation of human cells. The method, which involves the use of the antisense compound, is also useful for detecting the role of mdm2 expression in various cell functions and physiological processes and useful in both clinical research and
                                                                                                                                                                                                                Query Match
Best Local S
Matches 18
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modified_base
                 29-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of a
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                                              ABS67842;
                                                                                                                                                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleobases targeted a region (e.g. troof a nucleic acid encoding human mdm2.
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26-MAR-1999;
                                                                                                                                                                                                                                                                                                          oligonucleotides of
                                                                                                                                                                                                                                                                                                                        physiological processes and useful in both clinicadiagnostic tools. AAS29242-AAS29507 represent the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                An antisense compound, useful
                                                                          ABS67842
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MONIA B P.
COWSERT L M.
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                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                         TTTGTATTTTTAGTAGAGA 789
                                                                            standard;
                                                                                                                                                     TTTGTACTTTTAGTAGAGA 2
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                                                                                                                                                                                                                                                                             BP; 9 A; 4 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                  Conservative
               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nero P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98US-00048810
99US-00280805
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1. .20
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                                                                          DNA;
                                                                                                                                                                                                                                                                                                          the present invention
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                                                                            20
                                                                            BP
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                                                                                                                                                                                                                                  Pred.
                                                                                                                                                                                                                                               Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for treating e.g. cancer, comprises
(e.g. translation termination codon region)
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                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                .5e+03;
                                                                                                                                                                                                                                                DB 1;
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                                                                                                                                                                                                                                             Length 20;
                                                                                                                                                                                                                                                                                                                           human mdm2
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RESULT 1075 AAL40350 ID AAL4035

AAL40350 standard;

DNA;

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Human caspase 19-SEP-2002

6 antisense inhibition related oligo

SEQ ID

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69

(first entry)

Muscular; cytostatic; nootropic; neuroprotective; ophthalmological; antilipaemic; osteopathic; caspase 6; Rieger's syndrome; bone metabolism; ataxia telangiectasia; hyperproliferative disorder; cholesterol disorder;

Homo sapiens

apoptotic; haematopoietic

disorder;

cancer;

neurological;

disease

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                                                                                                                          Query Match
Best Local S
Matches 18
                                                                                                                                                                 for modulating the expression of human or mouse casein kinase 2-alpha prime. The antisense oligonucleotides are useful for inhibiting the expression of casein kinase 2-alpha prime, and for treating diseases or conditions associated with aberrant expression of casein kinase 2-alpha prime. Such diseases include diabetes mellitus, and hyperproliferative disorders (particularly cancers e.g. breader, prostate cancer, or liver cancer). The antisense compounds are also useful for diagnostics, therapeutics, prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits, and in distinguishing between functions of various members of a biological pathway. ABS67940-ABS67917 represent human or mouse casein kinase 2-alpha prime antisense oligonucleotides which comprise a phosphorothioate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; casein kinase 2-alpha prime; diabetes mellitus; hyperproliferative disorder; breast cancer; prostate cancer; liver cancer; infection; inflammation; tumour formation; cytosta antidiabetic; antiinflammatory; antimicrobial; phosphorothicate; antisense therapy; ss.
                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides targeted to nucleic acid encoding casein kinase 2-alpha prime, useful for diagnosing and/or treating a disease or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-627539/67.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-FEB-2001; 2001US-00780173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human casein kinase 2-alpha prime antisense
                                                                                                                                                         backbone
                                                                                                                                                                                                                                                                                                                                                                                          Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                                                       condition associated with expression of casein kinase 2-alpha
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                                                                                                                                                                                                                                                                                                                                                         present invention relates to
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20
                                                           l Similarity
18; Conserv
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                             CTCGGCTCACTGCAACCTC 987
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                                                                                                                                                                                                                                                                                                                                                                                          Page
CTCAGCTCACTGCAACCTC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Freier SM,
                                                                                                                       BP;
                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002WO-US002772
                                                                                                                                                                                                                                                                                                                                                                                          94; 129pp; English.
                                                                                                                         4.
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                                                                        1.8%;
94.7%;
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                                                             Pred. No. 1.5
); Mismatches
                                                                           Score 17.4;
Pred. No. 1.
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                                                                                                                         0 U;
                                                                           1.5e+03
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                                                                                          Length
                                                             Indels
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ion; cytostatic;
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ARESULT 1076
AAL40285/c
ID AAL4028
XX AAL4028
XX AAL4028
XX I9-SEP-
XX I9-SEP-
XX Muscula
KW AMISCULA
KW AITAILA
KW ATAXIA
KW ATAXIA
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KW ADOPTOO
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    in cells or tissues. The oligonucleotides can be administered therapeutically or prophylactically to treat an animal having a disease or condition associated with caspase 6, such as Rieger's syndrome or ataxia telangiettasia, hyperproliferative disorder, a haematopoietic disorder, a bone metabolism or cholesterol disorder, various types of cancer, neurological conditions such as Alzheimer's disease and other derepresents a human caspase 6 oligonucleotide relating to the invention.

NOTE: This phosphorothioate oligonucleotide sequence has 2'-MOE wings and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleotides in length that is targeted to a nucleic acid molecule encoding caspase 6, where the oligonucleotide specifically hybridises encoding caspase 6. Where the oligonucleotide of the with and inhibits the expression of caspase 6. The oligonucleotide of the invention specifically hybridises to and inhibits expression of caspase 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200229066-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-OCT-2000; 2000US-00679299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-OCT-2001; 2001WO-US030871.
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                                                                                                                                                                             antilipaemic; osteopathic; caspase 6; Rieger's syndrome; bone me ataxia telangiectasia; hyperproliferative disorder; cholesterol haematopoietic disorder; cancer; neurological; Alzheimer's disea
                                                                                                                                                                                                                                                       Caspase 6 antisense inhibition related PCR primer SEQ ID
                                                                                                                                                                                                                                                                                                                                                 AAL40285 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
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                                                                                                                                                                                                                       Muscular; cytostatic;
                                                                                                                                                                                                                                                                                      19-SEP-2002
                  04-OCT-2000;
                                                                                                                                                               apoptotic; human;
                                              03-OCT-2001; 2001WO-US030871.
                                                                              11-APR-2002
                                                                                                        WO200229066-A1
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                                                                                                                                    sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                           CTGGGATTACAGGTGTGAG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTGGGATTACAGGCGTGAG 883
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   caspase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                  2000US-00679299
                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           relates to an antisense oligonucleotide compound
n length that is targeted to a nucleic acid molecu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gonucleotide of 8 to 50 nucleotides in length 6, is useful for treating Rieger's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.8%;
94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 C;
                                                                                                                                                                                                                            nootropic; neuroprotective; ophthalmological;
                                                                                                                                                                 primer;
                                                                                                                                                                                                                                                                                                                                                 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8 G;
                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Watt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 17.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6 T;
                                                                                                                                                                  88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                          No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                            metabolism;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                disorder;
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RESULT 1077
AAL38206
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an antisense oligonucleotide compound of 8 to 50 nucleotides in length that is targeted to a nucleic acid molecule encoding caspase 6, where the oligonucleotide specifically hybridises with and inhibits the expression of caspase 6. The oligonucleotide of the invention specifically hybridises to and inhibits expression of caspase 6 in cells or tissues. The oligonucleotides can be administered therapeutically or prophylactically to treat an animal having a disease or condition associated with caspase 6, such as Rieger's syndrome or ataxia telangiectasia, hyperproliferative disorder, a haematopoietic disorder, a bone metabolism or cholesterol disorder, various types of cancer, neurological conditions such as Alzheimer's disease and other deregulated apoptotic pathological conditions. This polynucleotide sequence represents a human caspase 6 PCR primer relating to the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Brown-Driver VL,
                                                                                                                                                                                                                                                                                        haemostatic; BH3 interacting domain death agonist; liver disease; haematopoietic disorder; developmental disorder; immunological disorder; hyperproliferative disorder; apoptosis; human; chimeric; 2'-methoxyethyl
                                                                                                                 07-SEP-2000;
07-MAR-2001;
                                                                                                                                                        31-AUG-2001; 2001WO-US027316.
                                                                                                                                                                                  14-MAR-2002.
                                                                                                                                                                                                              WO200220547-A1.
                                                                                                                                                                                                                                         Chimeric
                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                            Human BH3 interacting domain death mRNA agonist inhibitor
                                                                                                                                                                                                                                                                                                                                                                                       29-AUG-2003
15-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAL38206 standard;
                                                                                       (ISIS-)
                                                                                                                                                                                                                                                                                                                                   Hepatotrophic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antisense oligonucleotide of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1006 GATTCTCCTGTCTCAGCCT 1024
                                       2002-393838/42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19
                                                                                         SISI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13; Page 85; 141pp;
                                                                                                                                                                                                                                                                              phosphorothicate backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             caspase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GATTCTCCTGCCTCAGCCT
                                                                Wyatt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                          PHARM INC
                                                                                                                  2000US-00657346.
2001US-00800631.
                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                  immunomodulatory;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; 3 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ucleotide of 8 to 50 nucleotides in length is useful for treating Rieger's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           μ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 17.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AT;
                                                                                                                                                                                                                                                                                                                                  cytostatic; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ŏ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20;
                                                                                                                                                                                                                                                                                                                                                                 SEQ
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                                                                                                                                                                                                                                                                                                                                                                 ID 49
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Novel antisense compound targeted to nucleic BH3 interacting domain death agonist, useful

acid molecule for treating a

e encoding the animals with

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RESULT 1078
AAL38189
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC targeted to a nucleic acid molecule encoding a BH3 interacting domain CC death agonist, where the compound specifically hybridises with and CC inhibits the expression of the BH3 interacting domain decompound of the invention is useful for inhibiting the expression of the BH3 interacting domain death agonist. The CC compound of the invention is useful for inhibiting the expression of the CC also useful for treating an animal having a disease or condition of associated with the BH3 interacting domain death agonist, e.g. CC haematopoietic disorder, hyperproliferative disorder, a developmental CC disorder, immunological disorder, or a disease or condition of the liver CC e.g., hepatitis, or a condition associated with apoptosis. The compound is useful for dispostics, therapeutics, prophylaxis and as research CC reagents and kits. This polymucleotide sequence represents an antisense C1 digonucleotide inhibitor of the DNA from human BH3 interacting domain CC death agonist RNA of the invention. NOTE: This sequence is a chimeric CC oligonucleotide 20 nucleotides in length, which is flanked on both sides by five-nucleotides. The internucleoside (backbone) linkages are CC phosphorothioate (P=5) throughout the oligonucleotide. (Updated on 29-AUG CC -2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 18
Novel antisense compound targeted to nucleic acid molecule encoding the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diseases associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                        Human BH3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-AUG-2003
15-AUG-2002
                                                                                                                                  07-SEP-2000; 2000US-00657346.
07-MAR-2001; 2001US-00800631.
                                                                                                                                                                                   31-AUG-2001; 2001WO-US027316.
                                                                                                                                                                                                                                                       WO200220547-A1
                                                                                                                                                                                                                                                                                                                                Hepatotrophic; immunomodulatory; cytostatic; antiinflammatory; hepatitis; haemostatic; BH3 interacting domain death agonist; liver disease; haematopoietic disorder; developmental disorder; immunological disorder; hyperproliferative disorder; apoptosis; human; chimeric; 2'-methoxyethyl; 2'-MOE; phosphorothioate backbone; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAL38189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAL38189 standard;
                                                                Zhang H,
                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                 2002-393838/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          729
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AGTAGCTGGGACTACAGGC 747
                                                                                                                                                                                                                                                                                                                                                                                                                                      interacting domain death mRNA agonist inhibitor SEQ
                                                                Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 6 A; 3 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                   PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               171pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BH3 interacting domain death agonist, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         멸
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 20
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N

GTTTCACCATGTTGGTCAG 20

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The invention relates to a compound 8 to 50 nucleotides in length CC targeted to a nucleic acid molecule encoding a BH3 interacting domain CC death agonist, where the compound specifically hybridises with and CC inhibits the expression of the BH3 interacting domain death agonist. The CC engound of the invention is useful for inhibiting the expression of the CC also useful for treating an animal having a disease or condition CC also useful for treating an animal having a disease or condition CC associated with the BH3 interacting domain death agonist, e.g. CC disorder, immunological disorder, or a disease or condition of the liver CC e.g., hepatitis, or a condition associated with appropriate The compound is useful for diagnostics, therapeutics, prophylaxis and as research CC e.g., hepatitis, or a condition associated with appropriate The compound CC is useful for diagnostics, therapeutics, prophylaxis and as research CC e.g., hepatitis, or a condition associated with appropriate The compound CC is useful for diagnostics, therapeutics, prophylaxis and as research CC oligonucleotide inhibitor of the DNA from human BH3 interacting domain CC death agonist RNA of the invention. NOTE: This sequence is a chimeric CC oligonucleotide of the invention. NOTE: This sequence is a chimeric CC oligonucleotide of the invention is lanked on both sides by five-nucleotides. The wings are composed of 2'-methoxyethyl (2' -MOE) nucleotides. The internucleoside (backbone) linkages are phosphorothicate (P=S) throughout the oligonucleotide. (Updated on 29-AUG cc 2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 3; Page 86; 171pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BH3 interacting domain death agonist, useful for treating animals with diseases associated with BH3 interacting domain death agonist, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hepatitis.
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Matches Query Match Best Local : Sequence 191 18; Similarity GTTTCTCCATGTTGGTCAG 209 20 BP; 3 A; 4 C; 6 G; 7 T; 0 U; 0 Other; Conservative 1.8%; 94.7%; 0; Score 17.4; DB 1; Pred. No. 1.5e+03; Mismatches ۳. Length Indels 0, Gaps 0

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RESULT 1079
AAD4294
XX
AC AAD4294
XX
AC AAD4294
XX
DT 15-NOV-
XX
Human P
XX
Human P
XX
Human P
XX
Homo sa
OS Synthet
FH modifie
FT modifie
                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                           modified_base
                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                antisense therapy;
                                                                                                                                                                                                                                                                           Human; antisense;
                                                                                                                                                                                                                                                                                                                        15-NOV-2002
                                                                                                                                                                                                                                                                                                                                              AAD42949;
                                                                                                                                                                                                                                                                                                                                                                   AAD42949 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                 PLA2,
                                                                                                                                                                                                                                                                                                group VI (Ca2+-independent) antisense oligo ISIS #129851
                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                               phospholipase A2; infection; inflammation;
; PLA2; phosphorothioate backbone; ss.
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modified\_base

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base=

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note= \*tag=

\_base= OTHER e= "2'-methoxyethyl (2'-MOE) nucleotides"

modified\_base

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base=

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base= OTHER == "2'-methoxyethyl (2'-MOE) nucleotides"

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"Phosphorothioate

backbone'

g)

modified\_base

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RESULT 1080
AAS96658/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to novel antisense compounds which inhibit the expression of phospholipase A2 (PLA2), group VI (Ca2+-independent). The invention is useful for inhibiting the expression of PLA2, group VI (Ca2+-independent) in human cells or tissues and for treating an animal, particularly a human suspected of having or being prone to a disease or condition associated with expression of human PLA2, group VI (Ca2+-independent). It is useful for diagnostics, therapeutics and as research reagent, e.g. prophylactically to prevent or delay infection, tumour formation or inflammation. The present DNA sequence is an antisense oligonucleotide targetted to human PLA2, group VI (Ca2+-independent) DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel antisense compounds useful for inhibiting gene expression of human phospholipase A2, group VI and for treating diseases associated with expression of phospholipase A2, group VI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-MAY-2001; 2001US-00851896
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bennett CF,
                                                                                                                                                                                   cell growth
                                                                                                                                                                                                 Telomerase
                                                                                                                                                                                                                      Telomerase reverse transcriptase, antisense oligonucleotide #68
                                                                                                                                                                                                                                                 09-APR-2002
                                                                                                                                                                                                                                                                                                  AAS96658 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC
                                                                                                                                  Synthetic
                                                                                                                                              Homo sapiens.
                      16-MAY-2000;
07-DEC-2000;
                                                         15-MAY-2001; 2001WO-US015774.
                                                                                                            WO200188198-A1
(-SISI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2002-616513/66
                                                                                                                                                                                                                                                                                                                                                                            851 GGCCTCCCAAAGTGCTGGG 869
                                                                                                                                                                                                                                                                                                                                                                                                               ll Similarity 94.7
18; Conservative
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ISIS
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                                                                                                                                                                                                                                                                                                                                                               GGTCTCCCAAAGTGCTGGG
                                                                                                                                                                                                reverse transcriptase; TERT; cytostatic;
                                                                                                                                                                                    inhibitor; antisense oligonucleotide; antisense technology;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Freier SM,
PHARM
                       2000US-00572423.
2000US-00733294.
                                                                                                                                                                                                                                                 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                           94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              5 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                   20
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                                                                                                                                                                                                                                                                                                                                                               20
                                                                                                                                                                                                                                                                                                                                                                                                                           Score 17.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                  apoptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                          20;
                                                                                                                                                                                                                                                                                                                                                                                                                 0
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The invention describes a compound, 8-50 nucleobases in length targeted contranscriptase), where the compound specifically hybridises with and contranscriptase), where the compound specifically hybridises with and contranscriptase), where the compound specifically hybridises with and contranscriptase of contranscriptase with and contranscriptase of contranscriptase were 20 nucleotides in length and composed of a central gap region consisting of the n 2'-deoxynucleotides, flanked on both sides (5' and 3' directions) by contranscriptase (RT)-polymerase chain reactions) by nucleotides. The wings were analysed for their effect on human contranscriptase (RT)-polymerase chain reaction (PCR). The compound is useful for inhibiting the expression of TERT in cells or tissues, for treating a human having disease or condition consociated with TERT, for modulating apoptosis, for inhibiting cell correction (prostics and therapeutics. This sequence is an antisense condition coligonucleotide used to modulate the activity of nucleic acid molecules encoding TERT, described in the method of the invention
RESULT 1081
ABS65070/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New compound targeted to nucleic acid molecule encoding telomerase transcriptase (TERT), which specifically hybridizes with and inhibits expression of TERT, useful for modulating apoptosis and inhibiting cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Monia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 5 A; 2 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 19; Page 91; 154pp; English
           modified_base
                                                                                                                  modified_base
                                                                                                                                                                     Key
modified_base
                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                          hyperproliferative
liver cancer.
                                                                                                                                                                                                                                                                                   ss; antisense;
                                                                                                                                                                                                                                                                                                             Human casein
                                                                                                                                                                                                                                                                                                                                        15-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                  ABS65070
                                                                                                                                                                                                                                                                                                                                                                                            ABS65070 standard; DNA; 20
                                                            modified_base
                                                                                                                                                                                                                                                                    cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1121 TCAAACTCCTGACCTCAGG 1139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002-075321/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ₿₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TCAAACTCCTGACCTCAAG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaarde WA,
                                                                                                                                                                                                                                                        antidiabetic; ant
erative disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                               kinase
                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                  casein kinase2-beta; human; itidiabetic; antiinflammatory;
            /note= "
16. .20
                       /mod_base=
/note= "2'-
                                                                                                                                                                                    Location/Qualifiers
                                                                                           /*tag= b
/mod_base=
                                                                                                                      1. .20
                                                    /*tag=
                                                                              note=
                                                                                                                                note=
                                                                                                                                                                                                                                                                                                               2-beta antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Freier
                                                                             se= OTHER
"Phophorothicate
                                                                                                                                    "A11
 Q.
                                                     a
                          -methoxyethyl residues
                                                                                                                                                                                                                                                                                                                                                                                               ΒP
                                        OTHER
                                                                                                                                                 OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>.</u>
                                                                                                                                 cytidines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>.</u>.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 17.4;
                                                                                                                                                                                                                                                         breast cancer; prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ľ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wancewicz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                      human; antisense gene therapy;
mmatory; diabetes; cancer; tumo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ٦;
                                                                                                                                    are
                                                                                backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                   5-methylcytidines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 20;
                                                                                                                                                                                                                                                                                                                #8
                                                                                                                                                                                                                                                            cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                          tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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RESULT 1082
ACC40949/c
ID ACC4094
XX
ACC ACC4094
XC ACC4094
XX
DT 23-MAY-
XX
DE Human s
XX
Human;
KW Human;
KW antiinf
KW hyperpr
KW ss.
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC The invention relates to a compound that is 8 - 50 nucleobases in length CC targeted to a nucleic acid molecule encoding Casein kinase 2-beta, and CC which specifically hybridises with and inhibits the expression of Casein CC which specifically hybridises with an 8-nucleobase CC portion of an active site on a nucleic acid molecule encoding Casein CC kinase 2-beta, and a carrier or diluent; (2) inhibiting the expression of Casein kinase 2-beta in cells or tissues by contacting the cells or tissues with the compound so that the expression of Casein kinase 2-beta in cells or tissues by contacting the cells or tissues with the compound so that the expression of Casein kinase 2-beta is inhibited; and (3) treating an animal having a disease or condition associated with casein kinase 2-beta by administering to the animal the cnew compounds of the expression of Casein kinase 2-beta is inhibited. The antisense compounds are useful for modulating the expression of Casein kinase 2-beta and for treating diseases or conditions associated with expression of Casein kinase 2-beta, e.g. diabetes or typerproliferative disorders, particularly cancer, such as breast cancer, configuration, therapeutics, prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and configuration transition between functions of various members of a biological pathway. The present sequence is an antisense oligonucleotide of the invention targeting human casein kinase 3-beta.
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 18
                                   Human; superoxide dismutase 1; antisense; neuroprotective; cytostatic; antiinflammatory; amyotrophic lateral sclerosis; apoptosis; hyperproliferative disorder; therapy; infection; inflammation; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotides targeted to nucleic acid encoding Casein kinase 2-beta, useful in diagnostic and research applications, or for treating a disease or condition associated with the expression of Casein kinase 2-beta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mckay R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-FEB-2001; 2001US-00780175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-JAN-2002; 2002WO-US003159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200262954-A2
                                                                                                                   Human superoxide dismutase 1 antisense inhibitor # ISIS 150503.
                                                                                                                                                                                                   ACC40949
                                                                                                                                                                                                                                         ACC40949 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; Page 91; 142pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-643409/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-AUG-2002
                                                                                                                                                          23-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                 989
                                                                                                                                                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                                                        18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention targeting human casein kinase 2-beta
                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                TCTGCCTCCCGGGTTCAAG 704
                                                                                                                                                                                                                                                                                                                                          TCTGCCTCCCAGGTTCAAG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Freier SM, Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 5 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "2'-methoxyethyl residues"
                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                      94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.8%;
                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                         ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                    <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 17.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                        .5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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RESULT 1083 AAL61497 ID AAL6149 XX

AAL61497

standard; DNA;

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                                                                                                                                          The invention relates to a compound of 8-50 nucleobases in length, CC targeted to a nucleic acid molecule encoding human superoxide dismutase 1. The compound specifically hybridises with and inhibits the expression of the compound specifically hybridises with and inhibits the expression of thuman superoxide dismutase 1 by hybridising with at least an 8-cc nucleobase portion of the nucleic acid molecule encoding the active site of the enzyme. The activity of compounds of the invention may be compounds of the invention may be compounds of the invention is antisense inhibition of human superoxide dismutase 1 expression by chimeric phosphorothicate coligonucleotides having 2'-methoxyethyl (2'-MOE) wings and a decay gap. CC compounds of the invention are useful for inhibiting the expression of human superoxide dismutase 1 in human cells or tissues, and for treating a disease or condition arising compounds of the invention are useful for inhibiting the expression of the invention associated with this enzyme (antisense therapy), cespecially amyotrophic lateral sclerosis, a disease or condition arising from aberrant apoptosis and a hyperproliferative disorder. It may also be used in diagnostics, therapputics and as a research reagent, e.g. compounds of the prevent or delay infection, inflammation or tumour condition. Sequences given in records ACC40880-ACC40957 represent human condition and antisense inhibitor oligonucleotides
                                                                     Matches
                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Key
modified_base
                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 15; Page 77; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel antisense compounds targeted to nucleic acids encoding human superoxide dismutase 1, for modulating expression of the dismutase and treating diseases or conditions, e.g. amyotrophic lateral sclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bennett FC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-184032/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-JUN-2002; 2002WO-US019664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003000707-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                     Local
                                  997
19
                                                                     1 Similarity
18; Conserv
                                    GGCTCAAGCGATTCTCCTG 1015
                                                                                                                                          20 BP; 6
                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001US-00888360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "Phosphorothioate linkages. All cytosines are
methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             *tag=
                                                                                                                                          A; 5 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                     94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ۵
                                                                     <u>.</u>
                                                                                       Pred. No.
                                                                                                    Score 17.4; DB 1;
                                                                     Mismatches
                                                                                       1.5e+03
                                                                                                      Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleotides'
                                                                       Indels
                                                                     0
                                                                     Gaps
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RESULT 1084
AAD47544/c
ID AAD4754
XX
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                                                                                                                Query Match
Best Local S
Matches 18
                                                                                                                                                                         for modulating the expression for activating transcription factor 3 (ATF3). ATF3 is also known as liver regeneration factor-1 (LRF-1), CRG-5, LRG-21, and TI-241. The invention is useful for the diagnosis, prevention and/or treatment of diseases or conditions associated with aberrant expression or activity of ATF3, such as ischaemia and diabetes. The antisense compound is useful in antisense therapy. The present sequence is an antisense oligonucleotide targetted to human ATF3 DNA. This sequence is used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                         New antisense oligonucleotide compounds, useful for preventing and/or treating conditions with aberrant activating transcription factor 3, such as ischemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAL61497;
                                                                                                                                                             Sequence 20 BP; 5 A;
                                                                                                                                                                                                                                                                                                      Example 15; Page 78; 126pp; English.
                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-441517/41.
                                                                                                                                                                                                                                                                                                                                                                                                   Baker BF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                08-NOV-2001; 2001US-00010002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-NOV-2002; 2002WO-US035331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003040161-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens
                                                                                          385
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATF3 antisense oligonucleotide, ISIS 185480.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                activating transcription factor 3; ATF3; ischaemia; diabetes;
                                                                                                                 18;
                                                                    _
                                                                                                                                                                                                                                                                                                                                                                                                                          ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          phosphorothicate backbone; antisense; ss.
                                                                                                                            Similarity
                                                                                         TCCCAAAGTGCTGGGATTA 403
                                                                                                                                                                                                                                                                                                                                                                                                   Dobie K;
                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag=
/mod_ba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "Phosphorothicate
methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'note= "2'methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    factor-1;
                                                                                                                          1.8%;
94.7%;
                                                                                                                                                             3 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "2'methoxyethyl
                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LRF-1; antisense therapy; CRG-5; LRG-21;
                                                                                                                 Pred. No. 1.56
); Mismatches
                                                                                                                                     Score 17.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   backbone;
                                                                                                                            .5e+03;
                                                                                                                                       DB 1;
                                                                                                                                       Length
                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A11
                                                                                                                                                                                                                                                                                                                                and diabetes.
                                                                                                                                                                                                                                                                                                                                           diagnosing, activity of the
                                                                                                                 0,
                                                                                                                  Gaps
                                                                                                                 0
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AAD47544 standard; DNA;

20 BP.

> Mus musculus Synthetic

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                                                                                                                                                                                                                                                       RESULT 1085
ADA20977
                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an Artemis nucleic acid coding for a protein involved in V(D)J recombination and/or DNA repair. Sequences of the invention are useful for treating severe combined immunodeficiencies (SCID) or cancer. They are also useful for diagnosing a patient, including a prenatal diagnosis with SCID, a predisposition to cancer, a immune deficiency or a carriage of a mutation increasing the risk of progeny to have such a disease. Peptides of the invention are used for preparing antibodies. The invention is useful in gene therapy. The present sequence is a PCR primer used to amplify human Artemis exon 6 [
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New ARTEMIS nucleic acid coding for a protein involved in V(D) \mbox{\it J} recombination and/or DNA repair, useful for treating and diagnosing severe combined immunodeficiencies (SCID) or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            severe combined
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAD47544;
                                                                 BCI2-associated X; BAX; nootropic; neuroprotective; antiparkinsonia anticonvulsant; ophthalmological; antidiabetic; virucide; antisense therapy; BAX antagonist; BAX inhibitor; familial amylotrophic lateral sclerosis; Alzheimer's disease; Parkinson's disease; Hodgkin's disease; cartilage-hair hyperplasia; diabetes-associated ocular disorder; scrapie infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                De Villartay J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-MAR-2001; 2001WO-IB000546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-MAR-2002; 2002WO-IB001737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200277026-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human;
                                                                                                                                                           Mouse BAX chimeric phosphorothioate oligonucleotide SEQ ID NO:150.
                                                                                                                                                                                                                                            ADA20977 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 68; 71pp; English.
                                                    aberrant apoptosis;
                                                                                                                                                                                        20-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                        778
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Artemis exon 6 amplifying PCR
                                                                                                                                                                                                                                                                                                               20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INSERM
                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                TTTTAGTAGAGATGGGGTT 796
                                                                                                                                                                                                                                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                               TTTTAGTGGAGATGGGGTT 2
                                                                                                                                                                                                                                                                                                                                                                                                                       ₿₽;
                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein; V(D)J recombination; DNA repair; gene therapy; d immunodeficiency; SCID; cancer; exon 6; PCR; primer; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INST NAT SANTE & RECH MEDICALE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Moshous D,
                                                                                                                                                                                                                                                                                                                                                                                                                         A,
                                                    mouse; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                1.8%;
94.7%;
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                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                          G; 3
                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                 Score 17.4;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                          Ή,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     primer,
                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                          ů,
                                                                                                                                                                                                                                                                                                                                                                                    .5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                          0 Other;
                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                     antiparkinsonian;
                                                                                                                                                                                                                                                                                                                                                                                                 20;
                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                      exon 6 DNA
                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                              δ
                                                                                                                                                                                                                                               The present invention describes a compound (I) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding BCL2-associated X (BAX) protein, where the compound specifically hybridises with the nucleic acid molecule encoding BAX protein and inhibits the expression of BAX protein. The compound specifically hybridises with at least 8-nucleobase portion of an active site on a nucleic acid molecule encoding BAX protein. Also described: (1) a composition comprising (I) and a pharmaceutical carrier or diluent; (2) inhibiting the expression of BAX protein in cells or tissues comprising contacting the expression of BAX protein (I); and (3) treating an animal having a disease or condition associated with BAX protein is inhibited. (I) has nootropic, neuroprotective, antiparkinsonian, anticonvulsant, ophthalmological, antidiabetic and virucide activities, and can be used in antisense therapy, and as a BAX antagonist. The antisense compounds (I) are useful for modulating the expression of BAX protein, e.g. familial amylotrophic lateral selections, Alzheimer's disease, Parkinson's disease, Hodgkin's disease,
                                                               Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compounds, useful for modulating the expression of BCL2-associated X (BAX) protein or for treating a disease or condition associated with BAX protein, e.g. Parkinson's disease, Hodgkin's disease or Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-JUL-2002; 2002WO-US022417
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modified_base
                                                                                                                                                                                 cartilage-hair hyperplasia, diabetes-associated ocular disorders or scrapte infection, or a condition that arises from aberrant apoptosis. The The compounds are useful as research reagents and in diagnostics. The present sequence represents a mouse BAX chimeric phosphorothicate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-JUL-2001; 2001US-00908147
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 3; Page 94; 139pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-239321/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
                                 394
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                                 GCTGGGATTACAGGCGTGC 412
                                                                                                                                   20
GCTGGGATTAAAGGCGTGC 19
                                                                 Conservative
                                                                                                                                   BP; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                                                                                                                                                  which is used in an example from
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/note= "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "phosphorothioate linkages, and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6. .20
                                                                                                                                   A.
                                                                                                                                   3 C; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            _base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "2'-0-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ი
                                                               ٥,
                                                                                                                                   G; 4 T; 0 U; 0 Other;
                                                                               Score 17.4;
Pred. No. 1.
                                                                 Mismatches
                                                                                 .5e+03
                                                                                                 DB 1;
                                                                                               Length 20;
                                                                                                                                                                    the
                                                                 Indels
                                                                                                                                                                    present invention
                                                               <u>,</u>
                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disease
                                                               0;
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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, IKBR, I-kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; ikappab r; antisense; immune response; infection; inflammation; tumour; prophylaxis; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighte immune response or infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human inhibitor-kappa B-R antisense oligonucleotide, ISIS #130450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAL61525 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                          Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-468635/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-NOV-2001; 2001US-00993731.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-NOV-2002; 2002WO-US035597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-MAY-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                       Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                        Page
                                                                                                                                                                                                                                                                                                                                                                                                                       Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry
                                                                                                                                                                                                                                                                       74; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mod_base= OTHER
/note= "Phosphorothicate backbone; All cytidine residues
re 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 *tag= c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 *tag= b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       _base= OTHEK
e= "2'-methoxyethyl {2'-MOE} nucleotides"
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                                                                                                                                                                                                                                                                                                                         a heightened
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NFKBIL2;
therapy;
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Query Match Best Local S Matches 18

Similarity

1.8%; 94.7%;

Score 17.4; Pred. No. 1. Mismatches

1.5e+03, DB 1;

Length Indels

0

Gaps

0

0;

18;

Sequence

20 BP; 4 A; 3 C; 9 G; 4 T; 0 U; 0 Other;

RESULT 1086 AAL61525

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RESULT 1087
ADD21684/c
ADD21691/
ID ADD2
XX
AC ADD2
XX
DT 15-J
XX
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                                                                                     RESULT 1088
                                                                                                                                                                           Matches
                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                           The invention comprises antisense oligonuclectides which are targeted to the human mdm2 gene. The antisense oligonuclectides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonuclectides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonuclectides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonuclectide of the invention. The present sequence contains 2'-methoxyethoxy-residues and has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human mdm2 antisense oligonucleotide #247.
                                                                                                                                                                                                                                                                                                                                                                                                                          Novel antisense compound targeted to 5' untranslated region, coding region, or intron:exon junction of nucleic acid molecule encoding mdm2, useful for treating e.g. cancer, psoriasis or restenosis by inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antisense oligonucleotide; human; mdm2; hyperprolife hyperproliferative disorder; cancer; psoriasis; fibratherosclerosis; restenosis; apoptosis modulation; p2'-methoxyethoxy-residue; phosphorothicate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADD21684 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Miraglia LJ,
Manoharan M;
                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                         Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-DEC-2001; 2001US-00005344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-DEC-2002; 2002WO-US038281.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS
             15-JAN-2004
                                                                                                                                                                                                                                                   phosphorothicate backbone.
                                       ADD21691;
                                                              ADD21691
                                                                                                                                                                                                                                                                                                                                                                                                                 expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2003-577263/54.
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                                                                                                                                                531
                                                                                                                          19
                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 249; 289pp; English
                                                                                                                          CATTCTCCTGCCTCAGCCT 1
                                                                                                                                                 CATCCTCCTGCCTCAGCCT
                                                                                                                                                                                                                          20
                                                              standard;
                                                                                                                                                                                                                          BP; 6
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                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PHARM INC
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               (first entry)
                                                                                                                                                                                                                          P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PS,
                                                              DNA;
                                                                                                                                                                                     1.8%;
                                                                                                                                                                                                                          2 C; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Graham MJ,
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                                                                                                                                                                          Score 17.4; D
Pred. No. 1.5e
0; Mismatches
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                                                                                                                                                                                                                          T; 0 U;
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                                                                                                                                                                                      1.5e+03;
                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                            0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Koller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        fibrosis;
                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           p21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'n
                                                                                                                                                                                                    20;
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RESULT 1089
ADD21692/c
ID ADD2169
XX ADD2169
XX ADD2169
XX IS-JAN-
XX Antiser
KW hyperp)
KW athero;
KW athero;
KW 2'-metl
XX
OS Homo s
XX
PM WO2003
XX
PD 12-JUN
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                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention comprises antisense oligonucleotides which are targeted to the human mdm2 gene. The antisense oligonucleotides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonucleotides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonucleotides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonucleotide of the invention. The present sequence contains 2'-methoxyethoxy-residues and has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense oligonucleotide; human; mdm2; hyperprolife hyperproliferative disorder; cancer; psoriasis; fib atherosclerosis; restenosis; apoptosis modulation; il 2'-methoxyethoxy-residue; phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel antisense compound targeted to 5' untranslated region, coding region, or intron:exon junction of nucleic acid molecule encoding mdm2, useful for treating e.g. cancer, psoriasis or restenosis by inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human mdm2 antisense oligonucleotide #254.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; SEQ ID NO 256; 289pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Miraglia LJ,
Manoharan M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-DEC-2001; 2001US-00005344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-DEC-2002;
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                                                                                         hyperproliferative disorder; cancer; psoriasis; fil atherosclerosis; restenosis; apoptosis modulation; 2'-methoxyethoxy-residue; phosphorothicate backbon
                                                                                                                                                               Human mdm2 antisense oligonucleotide #255.
                                                                                                                                                                                                                                                        ADD21692 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS
                                                              Homo sapiens
                                                                                                                                      antisense oligonucleotide; human; mdm2; hyperproliferation;
                                                                                                                                                                                              15-JAN-2004
                                                                                                                                                                                                                            ADD21692;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression.
                                                                                                                                                                                                                                                                                                                                                              578
                                                                                                                                                                                                                                                                                                                                    19
                                                                                                                                                                                                                                                                                                                                                                                             18;
                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                        CCACTACACCTGGCTAATT 596
                                                                                                                                                                                                                                                                                                                                    CCACCACACCTGGCTAATT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PS,
                                                                                                                                                                                                                                                                                                                                                                                                           1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        3 C; 8 G;
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                                                                                                                                                                                                                                                                                                                                                                                                             Score 17.4;
Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                         Length
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12-JUN-2003

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ADD71343/c
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention comprises antisense oligonuclectides which are targeted to the human mdm2 gene. The antisense oligonuclectides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonuclectides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonuclectides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonuclectide of the invention. The present sequence contains 2'-methoxyethoxy-residues and has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel antisense compound targeted to 5' untranslated region, coding region, or intron:exon junction of nucleic acid molecule encoding mu useful for treating e.g. cancer, psoriasis or restenosis by inhibit:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Miraglia LJ,
Manoharan M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GFAT 1 gene intron 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 9 A; 4 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-DEC-2002; 2002WO-US038281.
Judging relative onset risk of diabetes including type I or II diabetes and renopathy with or without type II diabetes accompanying, by detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                glutamine:fructose-6-phosphate amide transferase 1; ss; primer
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                                                                                      WPI; 2003-313261/30.
                                                                                                                                                                                                                                                              07-SEP-2001; 2001JP-00271870.
28-MAR-2002; 2002JP-00090861.
                                                                                                                                                                                                                                                                                                                                                  06-SEP-2002; 2002WO-JP009093
                                                                                                                                                                                                                                                                                                                                                                                                           20-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003023063-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diabetes; haplotype; polymorphism; diagnosis; renopathy; intron;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JAN-2004
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                                                                                                                                            Yasumo H,
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                                                                                                                                               Watanabe
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Pred. No. 1.
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by inhibiting
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The invention relates to a method of judging the onset risk of diabetes CC comprising detecting a haplotype consisting of gene polymorphism at 1 or CC more positions selected from (a)-(h) from a specimen containing human CC genomic DNA supplied by a patient: (a) the nucleotide located at position CC 36 of the intron 1 on GFAT1 (glutamine:fructose-6-phosphate amide CC transferase 1) gene (nucleotide number 62 in sequence ADD71329; (b) the CC nucleotide number 266 in sequence ADD71330; (c) the nucleotide located at position -147 of the intron 12 on GFAT1 gene (nucleotide number 338 in CC sequence ADD71331; (d) the nucleotide located at position 8 on GFAT1 gene (nucleotide located at position 8 on GFAT1 gene (nucleotide numbers 336-360 in sequence CC 12 on GFAT1 gene (nucleotide located at positions 1853-1877 of CC 12 on GFAT1 gene (nucleotide located at positions 1989-2007 of the intron CC 12 on GFAT1 gene (nucleotide numbers 328-347 in sequence ADD71333; (f) CC 12 on GFAT1 gene (nucleotide located at position 11 to -22 of the intron 18 on GFAT1 gene (nucleotide numbers 327-266 in sequence ADD71335; and (h) the nucleotide located at position 2632-2661 of the intron 3 on GFAT1 gene (nucleotide sequence ADD71335). The method is useful for judging relative onset risk of diabetes accompanying. This sequence represents a PCR primer used to amplify intron 8 of the GFAT1 gene in order to determine polymorphisms in the sequence of the GFAT1 gene in order to determine polymorphisms in the sequence of the GFAT1 gene in order to determine polymorphisms in the sequence of the GFAT1 gene in order to determine polymorphisms in the sequence of the seq
Sequence 20 BP; 8 A; 7 C; 1 G; 4 T; 0 U; 0 Other;
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                                                                           in the sequence.
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밁 S Query Match Best Local S Matches 18 778 20 18; Similarity TTTTAGTAGAGATGGGGTT 796 TTTTAGTAGAGACGGGGTT 2 Conservative 1.8%; ٥, Score 17.4; Pred. No. 1. Mismatches 1.5e+03; DB 1; Length Indels 20; <u>,</u> Gaps

0

RESULT 1091 ABZ99106 Human PDE4C oligonucleotide sequence. 17-OCT-2003 ABZ99106 standard; (first entry) DNA;

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; WO200285308-A2 Homo sapiens. inflammation; respiratory disease; ds.

31-OCT-2002.

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC.

Nyce JW, Miller S, Li Y, Sa, , Tang L, Sandrasagra A, Shahabuddin s Katz 'n Pabalan ŗ Aguilar D;

WPI; 2003-229219/22

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or

ubiquinone.

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RESULT 1092
ABZ97916
ID ABZ9791
XX
AC ABZ9791
XX
DT 17-OCT
DY 17-OCT
XX
Human;
KW Human;
KW antiis;
KW antiis;
KW antiis;
KW antiis;
KW antia;
KW lung
XX
PF 23-A
XX
PF 23-A
XX
PR (EP
XX
DR WP:
XX
WP:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the contribution codon, coding region, 5' or 3' end genomic flanking regions, coding and 3' intron-exon junctions, or regions within 2-10 nucleotides of coding a polypeptide associated with lung and/or coding a polypeptide associated with lung and/or coding a polypeptide associated with lung and/or coding an introduction and a second active agent comprising an coding an introduction and a second active agent comprising an coding antiinflammatory steroid and ubiquinone. A composition of the invention code in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also contiinflammatory steroid in a subject, for reducing or depleting levels of of or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing bronchoconstriction, coding surfactant in a subject, stissue, or treating bronchoconstriction, and subject, stissue, or treating bronchoconstriction, coding surfactant in a subject, stissue, or treating bronchoconstriction, coding surfactant in a subject, stissue, or treating bronchoconstriction, coding surfactant in a subject, stissue, or treating bronchoconstriction, coding surfactant in a subject, stissue, or treating bronchoconstriction, coding surfactant in a subject, stissue, or treating bronchoconstriction, coding surfactant in a subject, stissue, or treating bronchoconstriction, coding surfactant in a subject stissue, or treating bronchoconstriction, coding surfactant in a subject stissue, or treating bronchoconstriction, coding surfactant in a subject stissue, or treating bronchoconstriction, coding surfactant in a subject stissue, or treating bronchoconstriction, coding surfactant in a subject stissue, or treating bronchoconstriction, coding surfactant in a subject stissue, or treating bronchoconstriction, coding surfactant in a subject stis
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published not segmented.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 14348; 872pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchdilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human RANTES oligonucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                  WO200285308-A2
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                                                                                 Miller S,
                                                                                                        Мусе JW,
                                                                                                                                                                                                      24-APR-2001; 2001US-0286137P
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Tang L,
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                                                                                                     Sandrasagra
                                                                                                                                                           PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                 respiratory
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Shahabuddin
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Pred. No. 1
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                                                                                                        Katz E,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy;
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就是是
pharmaceutical composition for treating ailments associated with impaired
respiration, has oligo(s) antisense to specific gene(s) or its
corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                       SEQ
                                                                                                                                                       ID NO 13158; 872pp;
                                                                                                                                                       English
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cc nasal airway dysfunction and a second active agent comprising an cc antiinflammatory steroid and ubiquinone. A composition of the invention cc antiinflammatory steroid and ubiquinone. A composition of the invention common the invention common the invention cc immunosuppressive, and cytostatic activity. The composition may have a cc immunosuppressive, and cytostatic activity. The composition may have a cc preventing a respiratory, lung or malignant disease or condition, also cc for enhancing the prophylactic or therapeutic respiratory effect of an cc antiinflammatory steroid in a subject, for reducing or depleting levels of or reducing sensitivity to adenosine, reducing levels of adenosine cr creeptor, producing bronchodilation, increasing levels of ubiquinone or compute the sequence data for this patent is not represented in the printed confiction, the sequence data for this patent is not represented in the printed confiction, the print of the print The invention relates to a novel pharmaceutical composition, first active agent comprising an oligonucleotide antisense t initiation codon, coding region, 5' or 3' end genomic flanki junctions of genes encoding a polypeptide associated with lung and/or ftp.wipo.int/pub/published\_pct\_sequences intron-exon junctions, or regions within 2-10 nucleotides or 3' end genomic flanking which has the regions,

片 Query Match Best Local S Matches 18 Sequence 20 214 GTCTCGAACTCCCGACCTC 232 N 18; Similarity GTCTCGAACTCCTGACCTC 20 BP; 3 A; 8 C; 3 G; 6 T; 0 U; 0 Other; Conservative 94.7%; <u>,,</u> Score 17.4; Pred. Mismatches No. 1 .5e+03 DB 1; Length Indels 0 Gaps

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17-OCT-2003 ABZ98007; ABZ98007 standard; (first entry) DNA; 20

Human RANTES oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; inflammation; respiratory disease; ds

WO200285308-A2

31-OCT-2002

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

Nyce JW, Miller S, Li Y, Tang Sandrasagra A, ,, Shahabuddin S Katz 'n Pabalan Ģ Aguilar D;

WPI; 2003-229219/22

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or CC masal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine creceptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject stissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. CC lung inflammation, lung allergies, or a respiratory disease or condition. CC specification, but was obtained in electronic format directly from WIPO or a ferromated in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1094
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 18
                                                            Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 13249; 872pp; English.
                   WPI; 2003-229219/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-OCT-2003
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                                                                                                                                                                                                                                23-APR-2002; 2002WO-US013135
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                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABZ92731 standard;
                                                                                                                                                                              24-APR-2001; 2001US-0286137P
                                                                                                                                     (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                      inflammation;
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Tang L,
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                                                                                    Sandrasagra A,
                                                                                                                                                                                                                                                                                                                                                                                                                   respiratory disease; ds.
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94.7%;
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Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of beinginone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its responding RNAs, and glucocorticoid or non-glucocorticoid steroid or corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                     03-OCT-2002
                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                    primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                 Human; Artemis gene; DNA repair factor; metallo beta-lactamase; RS-SCID; chromosome 10; severe combined immunodeficiency; SCID1; cancer; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer used to amplify Human Artemis gene exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABV72400;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABV72400 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  junctions of genes encoding a polypeptide associated with lung and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel pharmaceutical composition,
                                                                                                                                                                                                        22-MAR-2001; 2001WO-IB000546.
                                                                                                                                                                                                                                              22-MAR-2001; 2001WO-IB000546.
                                                                                                                                                                                                                                                                                                                         WO200277228-A1
                                                                                                                                                                   (INRM ) INSERM INST NAT SANTE & RECH MEDICALE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.8%;
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Pred. No. 1.5e+03;
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Example 1; Page 65; 71pp; English

유

treating SCID

New isolated nucleic acid molecule of the Artemis gene, useful

De Villartay J,

Moshous D,

Fischer A;

PCR primers ABV72389-ABV72416 were used to amplify exons of the human Artemis gene. This gene encodes a V(D)J recombination and/or DNA repair factor that belongs to the metallo beta-lactamase superfamily, and whose mutations give rise to the human RS-SCID condition. The gene is localised to chromosome 10. The Artemis gene or its nucleic acid is useful for diagnosing or treating severe combined immunodeficiencies (SCIDs) or

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RESULT 1096
ABX14992/c
ID ABX1499
XX ABX1499
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Best Local S
Matches 18
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                                                                                                                                                                            New isolated nucleic acid molecule encoding a delta opioid receptor variant associated with an eating or energy homeostasis disorder, useful for diagnosing a genetic predisposition to such disorder, e.g. anorexia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                          11-MAY-2001; 2001US-0290016P
                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-MAY-2002; 2002WO-US014940
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    The invention relates to an isolated nucleic acid molecule encoding delta opioid receptor variant associated with an eating or energy homeostasis disorder. Also included are a delta opioid receptor varienceded by the nucleic acid, an isolated antibody that specifically
                                                                                                                                                                                                                                                                                                                                                              (BIOI-) BIOINVEST LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 778
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                                                                                                                  Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  opioid receptor OPRD1-1
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otide polymorphism; eating disorder; anorexia nervosa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first
                                                                                                                    19;
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/note= "A is cov
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chodamine (TAMRA) moiety"
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                                                                                                                  39pp;
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94.7%;
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                                                                                                                  English
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variant

The invention describes a method of selecting genetic markers as targets for nucleic acid sequence amplification comprising selecting each of the genetic markers according to a heterozygosity index of 0.5 or greater. Selecting and amplification of genetic markers are useful as targets for

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RESULT 1097
ACA88946/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cc recognises the delta opioid receptor variant, a vector comprising the nucleic acid, a host cell transformed to contain the vector, producing the polypeptide by culturing the host cell, identifying an agent which combinates the expression of the nucleic acid, diagnosing a genetic predisposition to an eating or energy homeostrasis disorder by detecting cc the presence or absence of the variant nucleic acid in a patient sample, cc an allele specific primer that detects a polymorphism in the gene cc encoding a delta opioid receptor associated with an eating or energy contain the variant nucleic acids. The variants are named OPRDI-1 to contain the variant nucleic acids. The variants are named OPRDI-1 to contain the variant nucleic acids. The variants are named OPRDI-1 to contain the variant opioid receptor gene is located on chromosome 1. The nucleic acid molecules and delta opioid receptor variant are useful for disorder, such as anorexia nervosa. The allele specific primer is useful contained with the disorder cited. The present sequence is a genotyping PCR probe for detecting the presence of a particular SNP (single nucleotide polymorphism) in a sample
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Best Local
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                                                                                                      Selecting genetic markers as targets for nucleic acid sequence amplification, useful for improving genetic testing, e.g. feta determination, comprises selecting each of the genetic markers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Selection and amplification of genetic markers PCR related
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                                                                 Claim 36;
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12-OCT-2001; 2001AU-00008235.
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determination;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1098
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                                                                                                         The invention relates to a novel isolated nucleic acid molecule comprising a variant gene associated with an eating disorder and selected from any of 119 polymorphisms with their corresponding genotyping in dataset, alleles and HGBASE identification, given in the specification. The novel nucleic acid molecule has polymorphisms in the serotonin receptor 1D, delta-opioid receptor, or dopamine receptor D2, which is useful in diagnostic and prognostic assays for eating disorders, in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bergen
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20-JUL-2001; 2001US-0306440P.
13-NOV-2001; 2001US-0331285P.
19-DEC-2001; 2001US-0340843P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               anorexia nervosa; bulimia nervosa; probe; ss.
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                                                                                                                                                                                                                                                                                                                                             Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PRIC-)
                                                                                                                                                                                                                                                                                                                                                                                                                         nucleic acid molecule having polymorphisms in the serotonin receptor delta-opioid receptor, or dopamine receptor D2, useful in diagnostic prognostic assays for eating disorders, such as anorexia and bulimia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2003-268122/26
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18; Conserv
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or 1D; delta-opioid receptor; dopamine receptor D2;
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                                                                                                                                                                                                                                                                                                                                             English.
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BP; 4 A; 3 C; 8 G;

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particular anorexia nervosa and

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nervosa. probe

polynucleotide invention

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RESULT 1099
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XX ABD2896
XX ABD2896
XX ABD2896
XX Human,
XW Human,
XW respira
XW respira
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                          CC comprising oligonuclectides, effective for alleviating
CC bronchoconstriction, respiratory tract inflammation, allergies and
CC reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors,
CC surfactant depletion or hyposecretion, when administered to a mammal. The
CC oligonucleotides are derived from a gene encoding or regulating
CC expression of a target polypeptide associated with lung airway or lung
CC dysfunction or cancer and can be anti-sense to the corresponding mRNA.
CC The invention also describes a kit, that comprises: (a) a delivery
CC device, in separate containers, (b) the oligonucleotides, (c)
CC instructions for adding a carrier and for use of the kit. The composition
CC of the invention has antiallergic, antiinflammatory, antiasthmatic,
CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a
CC beta-adrenergic agonist. The composition is useful for preventing or
CC treating a respiratory, lung or malignant disease. The administered
CC composition comprises oligo and is administered to reduce the production
CC or availability, or to increase the degradation of the target mRNA or to
CC pulmonary obstruction, and/or bronchoconstriction and/or lung
CC pulmonary obstruction and/or lung
CC pulmonary obstruction and/or lung
CC pulmonary obstruction and content and cont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes a novel composition
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1121
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Tang
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       allergies and/or surfactant
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L, Shahabuddin
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   hypoproduction
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RESULT 1100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                    Nyce J
Miller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
             comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine (A) or (A) recepto surfactant depletion or hyposecretion, when administered to a mammal. oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       pulmonary transplantation rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human
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                                                                                                                                                                                 nucleic acids associated bronchodilating agent.
                                                                                                                                                                                                                  Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to
                                                                                                                                                                                                                                                                   WPI; 2003-093058/08
                                                                                                                                                                                                                                                                                                                                                                                  24-APR-2001; 2001US-0286036P
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                                                                                                                                                   Claim
                                                                                                                  This invention describes a novel composition (a) a first active agent,
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Tang
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                                                                                                                                                   ID NO 13249;
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L, Shahabuddin
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Pred. No. 1
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                                                                   (A) receptors,
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                 or lung
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can be anti-sense

to the

corresponding mRNA.

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targe nucleic acids associated with lung airway or lung dysfunction,

targeted to

and

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The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, antiagesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition control of the reduce the amount of target polypeptide present in the lungs. The composition, allergies and/or bronchoconstriction and/or lung construction, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction.

CC distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system ce., in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system ce., in the target and the text the sum environment and thereby, to
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RESULT 1101
ABD30947
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Best Local
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                                                                                                   Nyce JW,
Miller S,
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                                                                   WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 17.4;
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                                                                                                     Katz
S;
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                                                                                                                     'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.5e+03;
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                                                                                                                         Pabalan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
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bronchodilating agent

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cc analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, andlargies, asthma, impeded respiration, respiratory hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system corresponding to the first any invaried affects dine to its sue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, reducing adenosine or humanerration. When administered to a mammal. The
                                                                                                                                                                                                     Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstrictio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20
                                                 31-OCT-2002.
                                                                                                                                                                                                                                                                                                                 Human PDE4C-derived oligonucleotide SEQ ID 14348.
                                                                                                                                                                                        respiratory distress syndrome; allergic rhinitis;
                                                                                                                                                                                                                                                                                                                                                       29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                            ABD32137 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GTCTCGAACTCCCGACCTC 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GTCTCGAACTCCTGACCTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID
                                                                                                                                                                          chronic obstructive pulmonary disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 3 A; 8 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NO 13158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.8%;
94.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 17.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 20;
                                                                                                                                                                                                             pulmonary vasoconstriction;
                                                                                                                                                                          pulmonary hypertension;
cancer; bronchitis;
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comprising oligonucleotides, effective for alleviating
comprising oligonucleotides, effective for alleviating
comprising adenosine sensitivity, levels of adenosine (A) or (A) receptors,
comprising adenosine sensitivity, levels of adenosine (A) or (A) receptors,
comprising adenosine sensitivity, levels of adenosine (A) or (A) receptors,
comprising of a target polyposecretion, when administered to a mammal. The
coligonucleotides are derived from a gene encoding or regulating
compression of a target polyposecretion, when administered to a mammal. The
coligonucleotides are derived from a gene encoding or regulating
compression of a target polypositide associated with lung airway or lung
comprises on reaccer and can be anti-sense to the corresponding mRNA.
comprises, in separate containers, (b) the oligonucleotides, (c)
composition for adding a carrier and for use of the kit. The composition
comprises of the invention has antiallergic, antiinflammatory, antiasthmatic,
composition for adding a carrier and for use of the kit. The composition
composition comprises oligo and is administered to reduce the production
composition comprises oligo and is administered to reduce the production
composition, allergies and/or bronchoconstriction and/or lung
composition, allergies and/or bronchoconstriction and/or lung
composition, allergies and/or bronchoconstriction are associated
composition, allergies, asthma, impeded respiration, respiratory
composition, allergies, asthma, impeded respiration, respiratory
composition, and/or surfactant hypoproduction are associated
composition, allergies, asthma, impeded respiration, respiratory
composition, and composition such as pulmonary vascoomstriction
composition and composition of the target polyposition and composition of the composition
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composition of target polyposition and composition of the composition
composition of target polyposition of the com
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Miller S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 15; SEQ ID NO 14348; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                               the oligonucleotides into products that free adenosine into e.g., lung, brain, heart, kidney, etc, tissue environment an prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-093058/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (EPIG-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention describes a novel composition (a) a first active
                                            lung, brain, he
nt any unwanted
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Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sandrasagra A,
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                                                                                                      the system
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S Best Loca Matches Query Match Local Similarity 352 18; N CTCCTGAGCTCAAGCAGTC 370 CTCCTGAGCTTAAGCAGTC Conservative 94.7%; 1.8%; 0 Pred. No. Score 17.4; DB 1; Length 20; Mismatches 1.5e+03; <u>.</u> Gaps

0

Sequence 20 BP; 5 A; 6 C; 4 G; 5 T; 0 U; 0 Other;

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RESULT 1103
ADF47745/c
ID ADF47745 standard; DI
XX
AC ADF47745;
XX
AC ADF47745;
XX
DT 26-FEB-2004 (first of the company of the
                                                                                                        human; 5-HT7 receptor promoter; barbiturate-inducible element; serotonin mediated response; gastrointestinal; neuroleptic; antidepressant; antimigraine; gene therapy; schizophrenia; depmigraine; affective disorder; sleep dysregulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene promoter related PCR primer.
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chromosome

depression;

23-APR-2002; 2002WO-US013143.

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RESULT 1104
ADH89041/C
ID ADH8904
XX ADH8904
AC ADH8904
XX ADH8904
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes an isolated nucleic acid molecule CC comprising: (a) nucleotides 1-3081 of the 3081-bp sequence of a human 5-CC HT7 receptor promoter region (see ADP47717), or a fragment exhibiting 5-CC HT7 receptor promoter activity; (b) the complementary strand of (a); or CC (c) a nucleic acid capable of hybridising under stringent conditions to CC (a) or (b). Also described: (1) an isolated regulatory element of the 5-CC (T) a nucleic acid capable of hybridising under stringent conditions to CC molecule; (3) a host cell transformed with the vector; (4) a method for identifying compounds which are modulators of human 5-HT7 receptor promoter enhancer activity; (6) a method for identifying compounds that modulate the activity of the barbiturate-inducible element CC within the 5-HT7 receptor promoter region; (7) a method for identifying CC compounds that modulate the activity of the barbiturate-inducible element CC within the 5-HT7 receptor promoter region; (8) a method for identifying CC compounds capable of modulating the 5-HT7 receptor promoter region; (8) a method for identifying CC compounds capable of modulating the 5-HT7 receptor promoter enhancer CC activity; and (9) a method for identifying polypeptides which bind to CC serotonin mediated responses. The 5-HT7 receptor promoter has CC gastrointestinal, neuroleptic, antidepressant and antimigraine CC activities, and can be used in gene therapy. The nucleic acid is useful in preparing a composition for treating conditions related to serotonin-mediated responses, e.g., schizophrenia, depression, migraine, affective buman 5 cd isorders selective.
                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New nucleic acid molecule exhibiting 5HT7 receptor promoter activity useful in preparing a composition for treating conditions related to serotonin-mediated responses, e.g., schizophrenia, depression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003102127-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-053452/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (JANC ) JANSSEN PHARM NV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-MAY-2003; 2003WO-EP005511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                       Human POLYX PCR primer #10
                                                               22-APR-2004
                                                                                                    ADH89041;
                                                                                                                                          ADH89041 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                         georders, sleep dysregulation or gastrointestinal functions. The human 5 greeptor promoter region is located on chromosome 10. The present squence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                   698
                                                                                                                                                                                                                                             19
                                                                                                                                                                                                                                                                                                                           18;
                                                                                                                                                                                                                                                                                 GATTACAGGCGTGAGCCAC 887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 32; 48pp; English
                                                                                                                                                                                                                                                                                                                                                                                                           BP; 3 A;
                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vanhoenacker PJP,
                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                           5 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                              94.7%;
                                                                                                                                                                                                                                                                                                                                                                   1.8%;
                                                                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                                                                Score 17.4; DB 1
Pred. No. 1.5e+03
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                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                   Length 20;
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compound that binds to the polypeptide in an amount sufficient to compound that binds to the polypeptide in an amount sufficient to compound that binds to the polypeptide in an amount sufficient to compound that binds to the polypeptide in an amount sufficient to compound that binds to the polypeptide in an amount sufficient to compound that the presence of or predisposition to a disease associated with altered levels of POLYX DNA or protein in a first mammalian subject, involving measuring the level of expression of DNA or the amount of C protein in a sample from the first mammalian subject amount of DNA or protein in a sample from a second mammalian subject cancunt of DNA or protein in a sample from a second mammalian subject can level of DNA or protein in the first subject can compared to the control sample indicates the presence of a secompared to the disease. The sequences of the invention are useful for treating or preventing a POLYX-associated disorder which involves cantibody is useful in the manufacture of a medicament for treating a syndrome associated with a human disease. This sequence represents a PCR primer used to amplify a human POLYX polynucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-SEP-1999;
20-SEP-1999;
13-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     encoding them. The invention also relates to an antibody that immunospecifically binds to a POLYX polypeptide, a method of determining the presence or amount of a POLYX polypeptide in a sample involving contacting the sample with a probe that binds to the polynucleotide and determining the presence or amount of the probe bound to the DNA, a method of identifying an agent that modulates the expression or activity of a POLYX polypeptide involving providing a cell expression the polypeptide, contacting the cell with the agent and determining whether the agent modulates expression or activity of the polypeptide indicates a alteration in expression or activity of the polypeptide indicates a modulation, and a method of modulating the activity of a polypeptide modulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; POLYX; PCR; ss; POLYX-associated disorder; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-OCT-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to human POLYX polypeptides and the polynucleotides encoding them. The invention also relates to an antibody that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shimkets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-SEP-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 5; SEQ ID NO 39; 93pp; English
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(HERR/)
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BOLDOG F L.
SMITHSON G.
RASTELLI L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FERNANDES E. HERRMANN J L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G RA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-0153629P.

99US-01547620P.

99US-0154762P.

99US-0159231P.

2000US-00659634.

2001US-0276960P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fernandes E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Herrmann JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Liu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ×
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19

1000 TCAAGCGATTCTCCTGTCT 1018

Query Match Best Local

18;

Conservative

0;

Mismatches

Indels

٥,

Gaps

0

.5e+03

DB 1; Length

Similarity

94.7%; 1.8%;

> Score 17.4; Pred. No. 1

Sequence

20

B₽;

7

A; 3 C; 7 G;

3 T;

0 U;

0 Other;

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RESULT 1105
ADJ59781
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                                                            Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                  The present invention relates to an oligonuclectide anti-sense to e.g., initiation codon, coding region with 2-10 nuclectides of 5'-end and 3'-end of nuclectic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonuclectide and optionally surfactant operatively linked to the oligonuclectide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonuclectide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            interleukin; II-4 receptor; II-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADJ59781 standard;
                                                                                                                           Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; SEQ ID NO 637; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-203534/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shahabuddin S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-JUL-2002; 2002US-0399076P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-JUL-2003; 2003WO-US023509
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                      (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airwobstruction. The present sequence represents an oligonucleotide of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (EPIG-) EPIGENESIS PHARM INC
                               214
N
                                                                             Similarity
                             GTCTCGAACTCCCGACCTC 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tang L,
CTCGAACTCCTGACCTC 20
                                                              Conservative
                                                                                                                         BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             associated
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                                                                           1.8%;
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                                                                                                                         8
                                                                                                                         C; 3
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                                                            Score 17.4; D
Pred. No. 1.5e
0; Mismatches
                                                                                                                         G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             #30.
                                                                             .5e+03
                                                                                          DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Miller
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de of the
                                                            Gaps
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RESULT 1106 ADJ60991 ID ADJ6099

ADJ60991 standard; DNA; 20 BP

RESULT 1107 ADJ59872 ID ADJ5987

ADJ59872 standard;

DNA;

20 BP

ADJ59872;

06-MAY-2004

(first entry)

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                                                                                                                                                                                        The present invention relates to an oligonucleotide anti-sense to e.g., cc initiation codon, coding region with 2-10 nucleotides of 5' end and 3'-ce end of nucleic acid target comprising gene (s) chosen from e.g. cc interleukin (IL)-4 receptor, IL-5 receptor or salts of the cc oligonucleotide and optionally surfactant operatively linked to the cc oligonucleotide. The method is useful for preventing or treating a CC respiratory or lung disease, which involves administering to the airways cf a subject an effective amount of an inhibitor. The oligonucleotide is cc useful for production of a medicament for the prevention and/or treatment cc useful for production of a medicament for the prevention and/or treatment cc respiratory or lung disease. The respiratory or lung disease is cc chosen from airway inflammation, allergy(ies), asthma, impeded cc respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway construction. The present sequence represents an oligonucleotide of the
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                                                                      Matches
                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nyce JW, Tang L,
Shahabuddin S, Lu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligonucleotide associated to PDE4C #57
                                                                                                                                             Sequence 20 BP; 5 A; 6 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-203534/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-JUL-2002; 2002US-0399076P
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                                   352
                                                                      18;
N
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                                   CTCCTGAGCTCAAGCAGTC 370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 1847; 85pp; English.
CTCCTGAGCTTAAGCAGTC 20
                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sandrasagra A,
                                                                                       1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cong H;
                                                                      ٥,
                                                                                           Pred. No.
                                                                                                          Score 17.4; DB 1;
                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Aguilar
                                                                                           1.5e+03
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                                                                                                          Length 20;
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                                                                      Gaps
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RESULT 1108
ADK43371/c
ID ADK4337
XX ADK4337
AC ADK4337
XZ
DT 06-MAY-
XX
DE Human F
XX
CW PTPRA;
KW LCA-rel
KW hyperpx
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                                                                                                                                                                                                                                                                                                                                á
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-cend of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the coligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is cohesen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway character and the collection of the prevention and or treatment of the prevention and or treatment of comparison of the prevention and/or treatment of a respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway
                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        interleukin; IL-4 receptor; IL-5 receptor; lung disease;
airway inflammation; allergy; asthma; impeded respiration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2004011613-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shahabuddin
                                                                                                                                                                                             371/c
ADK43371 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (EPIG-) EPIGENESIS PHARM INC
PTPRA; protein tyrosine phosphatase, receptor type alpha; LCA-related phosphatase; LRP; HLPR; HPTPA; PTPRL2; RPTPA; cytostatic hyperproliferative disorder; metabolic; antisense target; human; ds
                                                                                                                      06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             obstruction. The present sequence represents an oligonucleotide
                                                                               Human PTPRA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sapiens
                                                                                                                                                                                                                                                                                                                                    728
                                                                                                                                                                                                                                                                                                                                                                        1 Similarity
18; Conserv
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                                                                                                                                                                                                                                                                                                                                  GAGTAGCTGGGACTACAGG 746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 728; 85pp; English.
                                                                                                                                                                                                                                                                                             GAGTAGCTGGGATTACAGG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tang L, Sai
in S, Lu H,
                                                                               DNA targeted for antisense therapy -
                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sandrasagra
H, Cong H;
                                                                                                                                                                                                                                                                                                                                                                                         1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                 20 BP
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                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                             Score 17.4; DB 1;
Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RANTES #121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Þ,
                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Aguilar D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Miller
                                                                                                                                                                                                                                                                                                                                                                                                                 Length 20;
                                                                                 SEQ ID 195
                                                                                                                                                                                                                                                                                                                                                                            Indels
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                        cytostatic;
                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     õ
                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel compound 8-80 nucleobases in length which is targeted to and specifically hybridises with a nucleic acid molecule encoding PTPRA (protein tyrosine phosphatase, receptor type alpha, LCA-related phosphatase; LRP; HLPR; HPTPA; PTPRL2; RPTPA) and inhibits the expression of PTPRA. The compound of the invention demonstrates cytostatic activities and may be useful for treating a disease or condition associated with PTPRA, such as a hyperproliferative disorder or metabolic disorder as well as in research and disgnostics for modulating the expression of PTPRA. The current sequence is that of a human PTPRA DNA of the invention which was targeted for antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2004011623-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding protein tyrosine phosphatase receptor type alpha (PTPRA), useful for treating hyperproliferative or metabolic disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-JUL-2002; 2002US-00210556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-JUL-2003; 2003WO-US023972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 16; SEQ ID NO 195; 289pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-143851/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ISIS-)
                                                                                                                 modified_base
                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                       PTPRA; protein tyrosine phosphatase; receptor type alpha; LCA-related phosphatase; LRP; HLPR; HPTPA; PTPRL2; RPTPA; cytos hyperproliferative disorder; metabolic; antisense; ss; human; hyperproliferative disorder; metabolic; antisense; ss; human; 2'-MOE wing; 2'-methoxyethyl gapmer; phosphorothioate backbone.
                                                                                                                                                                                                                                                              Antisense 2'-MOE gapmer oligo targeted to human PTPRA -
                                                                                                                                                                                                                                                                                                                                                        ADK43253 standard; DNA;
               WO2004011623-A2
                                                                                                                                                                                                                                                                                               06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                    20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20
                                                                                                                                                                                                                                                                                                                                                                                                                                    CAGGCTGGTTTCGAACTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Freier SM,
                                                                                                                                                                                                                                                                                               (first entry)
                                           /mod_base= OTHER
/mod_base= OTHER = Bases 1-5 and 16-20 comprise 2'-
/mote= "OTHER = Bases 1-5 and 16-20 comprise 2'-
methoxyethyl (2'-MOE) wings. Phosphorothicate backbone
throughout. All cytidines are 5-methylcytidines"
                                         throughout.
                                                                                                 /*tag=
                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dobie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 17.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The current sequence is that of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                       alpha;
RPTPA; cytostatic;
                                                                                                                                                                                                                                                                    SEQ
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RESULT 1110
ADJ10489
ID ADJ1048
XX ADJ1048
XX ADJ1048
XX Phospho
XX Phospho
XX Phospho
XX Phoman;
KW PpMT; P
KW PPMT; P
KW PPMT; P
KW Cardiov
KW ICMT; a
XX Homo sa
OS Synthet
XX Key
FT modifie
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FT modifie
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a novel compound 8-80 nucleobases in length which is targeted to and specifically hybridises with a nucleic acid molecule encoding PTPRA (protein tyrosine phosphatase, receptor type alpha, LCA-related phosphatase; LRP; HLPR; HPTPA; PTPRL2; RPTPA) and inhibits the expression of PTPRA. The compound of the invention demonstrates cytostatic activities and may be useful for treating a disease or condition associated with PTPRA, such as a hyperproliferative disorder or metabolic disorder, as well as in research and dispnostics for modulating the expression of PTPRA. The current sequence is that of an antisense 2'-MOE (2'-methoxyethy!) gagmer oligonucleotide which was targeted to human PTPRA of the invention.
                                                                                                                                                                                                                                                                                                   human; isoprenylcysteine carboxyl methyltransferase; ss; PCCMT; pcMTase; PPMT; PPMTase; HSTE14; MST098; MSTP098; growth factor signal transduction; cell replication; vesicular transport;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New compounds, particularly antisense oligonucleotides targeted nucleic acid encoding protein tyrosine phosphatase receptor type (PTPRA), useful for treating hyperproliferative or metabolic dis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-143851/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-JUL-2002; 2002US-00210556.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADJ10489 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 15; SEQ ID NO 77; 289pp; English.
              modified_base
                                                                                           modified_base
                                                                                                                                                        modified_base
                                                                                                                                                                                                                                                     hyperproliferative disorder; cancer; inflammatory; hypertension; cardiovascular; cytostatic; antiinflammatory; hypotensive; cardicum; antisense; phosphorothioate backbone; 2' MOE wing.
                                                                                                                                                                                                                                                                                                                                                                    Phosphorothicate antisense DNA oligo to modulate human ICMT SeqID 16
                                                                                                                                                                                                                                                                                                                                                                                                   17-JUN-2004
                                                                                                                                                                                                                         sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           207 CAGGCTGGTCTCGAACTCC 225
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CAGGCTGGTTTCGAACTCC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 3 A; 6 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                       /*tag= a
/mod_base= OTHER
/note= "OTHER= 2'
                               cytidine
                                                                                                                             /mod
                                                                                                                                                                          Location/Qualifiers
                                                                                                           'note=
                  6. .20
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                                                                                                                             base= OTHER
                                                                                                           "OTHER= phosphorothicate backbone"
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                                 nucleobases
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Pred. No. 1.5e+03;
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                               methoxyethyl (2' MOE) nucleotides.
ses are 5-methylcytidine."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   eted to a
type alpha
c disorder.
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RESULT 1111
ADJ10565/c
ID ADJ1056
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention relates to a novel antisense compounds that modulate the expression of isoprenylcysteine carboxyl methyltransferase (also known as ICMT, pcMTase, PPMT, ppMTase, HSTE14, MST098 and MSTP098) and CC located on chromosome 1p36. Specifically, it refers to compositions CC useful for inhibiting the expression of isoprenylcysteine carboxyl CC methyltransferase, which normally participates in cellular events such as CC growth factor signal transduction, cell replication, vesicular transport and the post-translational modification of the Ras family of GTPases. The CC present invention describes antisense oligonucleotides that comprise at CC least one modified nucleobase, a 5-methylcytosine. Accordingly, these compounds are useful for treating a disease or condition associated with isoprenylcysteine carboxyl methyltransferase such as a hyperproliferative CC disorder (e.g. cancer), an inflammatory condition, hypertension or CC cardiovascular disease. As such, they exhibit cytostatic, antisense only and are useful for research reagents and in diagnostics. This oligonucleotide sequence is a phosphorothicate antisense DNA oligo used to modulate human and carboard mathyline and carboard mathyline and carboard mathyline of the compounds of the compounds.
Matches
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Best Local
           human; isoprenylcysteine carboxyl methyltransferase; ss; PCCMT; pcMTase; ppMT; ppMTase; HSTE14; MST098; MST098; growth factor signal transduction; cell replication; vesicular transport; hyperproliferative disorder; cancer; inflammatory; hypertension; cardiovascular; cytostatic; antiinflammatory; hypotensive; cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 7 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     isoprenylcysteine carboxyl methyltransferase expression in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New compounds, particularly antisense oligonucleotides targeted to nucleic acid encoding isoprenylcysteine carboxyl methyltransferase, useful for treating cancer, hypertension, or cardiovascular or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-081071/08
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                                                                                                                                                           17-JUN-2004
                                                                                                                                                                                             ADJ10565;
                                                                                                                                                                                                                                 ADJ10565 standard; DNA;
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                                                                                                                      Target DNA
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                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                               1000 TCAAGCGATTCTCCTGTCT 1018
                                                                                                                                                                                                                                                                                                                                                                                              18;
                                                                                                                                                                                                                                                                                                                         N
                                                                                                                                                                                                                                                                                                                       TCAAGCGATTCTCCTGCCT 20
                                                                                                                      oligo
                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disease.
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                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= OTHER
/note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides.
cytidine nucleobases are 5-methylcytidine."
                                                                                                                        for
                                                                                                                                                                                                                                                                                                                                                                                                             94.7%;
                                                                                                                    antisense therapy of human
                                                                                                                                                                                                                                                                                                                                                                                                                                  1.8%;
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Pred. No. 1.5e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
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                                                                                                                         ICMT
                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
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                                                                                                                        SeqID
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This invention relates to a novel antisense compounds that modulate the CC expression of isoprenylcysteine carboxyl methyltransferase (also known as CC ICMT, PCCMT, pcMTase, PPMT, PPMTase, HSTE14, MST098 and MSTP098) and CC located on chromosome 1p36. Specifically, it refers to compositions CC useful for inhibiting the expression of isoprenylcysteine carboxyl CC methyltransferase, which normally participates in cellular events such as CC growth factor signal transduction, cell replication, vesicular transport CC and the post-translational modification of the Ras family of GTPases. The CC least one modified sugar moiety, a 2'-O-methoxyethyl (2' MOE) and at CC least one modified nucleobase, a 5-methylcytosine. Accordingly, these CC compounds are useful for treating a disease or condition associated with isoprenylcysteine carboxyl methyltransferase such as a hyperproliferative CC disorder (e.g. cancer), an inflammatory condition, hypertension or CC cardiovascular disease. As such, they exhibit cytostatic, antiinflammatory, hypotensive and cardiant activities and are useful for research reagents and in diagnostics. This oligonucleotide sequence is a CC DNA oligo representing a preferred target site for antisense therapy in human isoprenylcysteine carboxyl methyltransferase, given in an CC exemplification of the invention.
                                                                                                                                                                                                                                                                   RESULT 1112
ADM13970/c
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New compounds, particularly antisense oligonucleotides targeted to nucleic acid encoding isoprenylcysteine carboxyl methyltransferase, useful for treating cancer, hypertension, or cardiovascular or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-MAY-2002; 2002US-00159834.
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                                                    chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory;
                                                                                                                                                                              01-JUL-2004
                                                                                                                                                                                                                                                      ADM13970 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS
               neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                             Human mPGES-1
                                                                                                                                                                                                                                                                                                                                                                                 1000
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                                                                                                                                                                                                                                                                                                                                                                                                                      18;
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                                                                                                                                                                                                                                                                                                                                                                               TCAAGCGATTCTCCTGTCT 1018
                                                                                                                                                                                                                                                                                                                                               TCAAGCGATTCTCCTGCCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 92; 62pp;
                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disease
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                                                                                                                                                                              (first entry)
                                                                                                                                             chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; 3 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                      1.8%;
   arthritis; diabetes; cancer; ischaemia;
                                                                                                                                             antisense
                                                                                                                                                                                                                                                      20
                                                                                                                                                                                                                                                      ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 17.4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English
                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                            oligonucleotide SEQ ID NO:157.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                  The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpqES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to Q934.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC mpGES-1 in cells or tissues; and (3) a method of the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisfiammatory, neuroprotective, and antisense compounds have cytostatic, cophthalmological, immunomodulator, cardiant, neuroprotective, cophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or CC condition associated with mpGES-1-e.g., inflammation, Alzheimer's CC ophthalmoic, immunological, cardiovascular or neurological disorder.
                                                     Matches
                                                                   Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-SEP-2002;
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                                                                                                          Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; SEQ ID NO 157; 132pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                              lschemia.
                                                                 Local
                         390
20
                                                     18;
                                                                    Similarity
                            AAGTGCTGGGATTACAGGC 408
                                                                                                             BP; 3 A;
                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2002US-0413549P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
 GGGATGACAGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "phosphorothioate linkages and residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    . 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               _base=
                                                                                                             8 C; 3 G; 6 T; 0 U;
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                                                                 94.7%;
                                                                                1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "2'-O-methocyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER
                                                     0; Mismatches
N
                                                                   Score 17.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                   English.
                                                                    .5e+03
                                                                                                              0 Other;
                                                                                  DB 1;
                                                                                Length
                                                        Indels
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                                                        0
                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            e.g.,
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The present sequence represents a chimeric antisense oligonucleotide chargeted to human microsomal prostaglandin E2 synthase (mGCES-1). The human mGGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, chargeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding chibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal chargeted to a compound the compound conting a disease or condition associated with mPGES-1. MPGES-1 chimeric charing a disease or condition and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic, antistinflammatory, neuroprotective, notropic, antiarthritic, vasotropic, antiinflammatory, neuroprotective, notropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nottropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
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                                                                                                                                                                                                                                                                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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16. .20
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residues are 5-methylcytidines"
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"2'-0-methoxyethyls"
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RESULT 1114
ADM15339/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                            25-SEP-2002; 2002US-0413549P
                                                                                                                                                                                                                                                                                                                                                                                                25-SEP-2003; 2003WO-US030374
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/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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1. .20
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note= "2'-O-methocyethyls"
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94.7%;
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Pred. No. 1.5e+03;
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Claim 4; SEQ ID NO 1526; 132pp; English

The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The

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RESULT 1115
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Best Local S
Matches 18
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1 Similarity 94.7%;
18; Conservation
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residues are 5-methylcytidines"
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                                                                                                "2'-O-methoxyethyls"
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Pred. No. 1.
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RESULT 1116
ADM14492/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4; SEQ ID NO 901; 132pp; English
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                                                                                                                                                                                                                                                          microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; immunomodulator; cardiant; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene thorapy; inflammation; immunomodulatory; cardiovascular; gene thorapy; inflammation;
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                                                                                                                                                                                                             Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disor
                                                                                                                                                                                                                                                                                                                                                                         chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibi
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human mPGES-1 chimeric antisense oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADM14492;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADM14492 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (PHAA ) PHARMACIA CORP
                                                                                                         Synthetic
                                                                                                                                                                                       cardiovascular
                                                                                                                                  sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGGGTTCAAGCGATTCTCC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CGGGCTCAAGCGATTCTCC 1013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 5 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
                                                                                                                                                                                       disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94.7%;
                                                                                                                                                                                       neurological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 17.4; DB 1
Pred. No. 1.5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
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                                                                                                                                                                                                                      disorder;
                                                                                                                                                                                                                                                                                                                                                                         inhibitor;
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Gaps

0

modified\_base

Location/Qualifiers
1. .20
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RESULT 1117
ADM14702/c
ID ADM1470
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AC ADM1470
AC ADM1470
XX
AC ADM1470
XX
DT 01-JUL-
XX
DE Human n
XX
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                                                                                                                                                                                                                                                                                                                                                                  The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically horidise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisting lammatory, neuroprotective, nouroprotective, antistinflammatory, neuroprotective, nootropic, antistinflammatory, neuroprotective, nootropic, antistinglandialpological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                      Query Match
Best Local :
                   Human mPGES-1 chimeric antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 9
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                                                                                                                     ADM14702 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                    Local
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                                                  (first entry)
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                        A; 2 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                      1.8%;
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e= "2'-O-met
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                                                                                                                                                                                                                                                                                      Score 17.4; DB 1;
Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                       DB 1;
                   SEQ
                                                                                                                                                                                                                                                                                                     Length 20
                                                                                                                                                                                                                                                                         Indels
                   ID NO:889
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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin R2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin R2 synthase; mpGES-1; mpGES-1 inhibitor; antidiabetic; inmunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                     antisense oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic, antisinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues, and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
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modified_base
Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4; SEQ ID NO 889; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PHAA ) PHARMACIA CORP
  BP;
12 A; 2 C; 1 G; 5 T; 0 U; 0 Other;
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/note= "phosphorothioate linkages and
residues are 5-methylcytidines"
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/note= "2'-O-methocyethyls"
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Query Match Best Local Similarity Matches 18; Conserv

Conservative

0

Score 17.4; DB 1; Pred. No. 1.5e+03; 0; Mismatches 1

DB 1;

Length Indels

0;

Gaps

0

1.8%;

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RESULT 1118
ADM14961/c
ID ADM1496
XX ADM1496
XX ADM1496
XX Chimeri
KW Chimeri
KW microso
KW mocroso
KW microso
KW mocroso
KW mocroso
FT modifie
FT Tolaim
XX New mp1; 20
XX New mp1; 20
XX New mp1; 20
XX New microso
FX Claim x
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory;
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                     The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of
                                                                                                                                                                                                                                                                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-305094/28.
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                                                                                                                                                                                                                                                                  Claim 4; SEQ ID NO 1148; 132pp; English
      mPGES-1 in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1065
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      cells or tissues; and (3) a method of treating an animal
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e= "2'-O-methoxyethyls"
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inhibitor
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RESULT 1119
ADM15080/c
ID ADM1508
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric; antisense oligonucleotide; phosphorothioate;
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                            Gierse
                                                                                                                  25-SEP-2003; 2003WO-US030374
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                                                                                     25-SEP-2002; 2002US-0413549P
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/mod_base= OTHER
/note= "2'-O-methocyethyls"
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/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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/mod_base=
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Pred. No. 1
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inhibitor;
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WPI; 2004-305094/28

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Best Local S
Matches 18
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               modified_base
                                                                                                                                                                                                                                                                                                                  microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
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                                                                                                                                                                            modified_base
                                                                                    modified_base
                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric; antisense oligonucleotide; phosphorothioate; human;
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               /note= "
                                                                                                       residues
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residues are 5-methylcytidines"
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                                 "2'-O-methocyethyls
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Pred. No. 1
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RESULT 1121
ADM14687/c
ID ADM1468
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XX DE Human m
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KW microsc
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Best Local Similarity
Matches 18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, noutropic, antiinflammatory, neuroprotective, nootropic, antiinflammatory, neuroprotective, nootropic, antiinflammatory, neuroprotective, nootropic, antiinflammatory, and cardiovascular activities, and can obe used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                             chimeric; antisense oligonucleotide; phosphorothioate; human; microseomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microseomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
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                    Alzheimer's disease; arthritis; diabetes; cancer; isch
reperfusion injury; ophthalmic disorder; immunological
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                                                                                                                                                                                                                                                                                                                                                         standard;
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'note= "2'-O-methoxyethyls"
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Pred. No. 1.
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                                                cancer; ischaemia;
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RESULT 1122
ADM13931/c
ID ADM13931 standard; DNA; 20 BP
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense compound, having a sequence targeted to a nucleic acencoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-305094/28
                                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                       antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; SEQ ID NO 874; 132pp; English
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The CC thuman mpGES-1 gene is located on chromosome 9, more specifically to CC 9q34.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, antiinflammatory, neuroprotective, nootropic, antisense oligonucleotides and in gene therapy. The antisense compound CC condition associated with mpGES-1 inclammation, Alzheimer's CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC condition associated with mpGES-1 e.g., inflammation of the condition inflammation inflammatical condition inflammation inflammatical condition inflammat
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/note= "2'-O-methocyethyls"
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SXS
RESULT 1123
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                    Claim 4; SEQ ID NO 1614; 132pp; English.
                                                       New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.c inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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/mod_base= OTHER
/mote= "2'.-O-methocyethyls"
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/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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                                                         treating e.g.,
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present sequence represents a chimeric antisense oligonucleotide

25-SEP-2003; 2003WO-US030374

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RESULT 1124
ADM14038/c
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Matches 18
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Synthetic.
                                                                                                                                                                                            modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 8 A; 6 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antisense oligonucleotides and antisense compounds have cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              targeted to human microsomal prostaglandin E2 synthase (mPGES-1).
 08-APR-2004.
                                    WO2004028458-A2
                                                                                                                        modified_base
                                                                                                                                                                                                                                                                                        modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADM14038;
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                                                                                                                                                                                                                                                                                              Location/Qualifiers
1. .20
                                                                                                                                                                                                                   /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                            16.
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/note= "2'-O-methocyethyls"
                                                                                                          *tag=
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                                                                        base= OTHER
== "2'-O-methoxyethyls"
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Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The CC human mPGES-1 gene is located on chromosome 9, more specifically to Q334.3. The present invention also describes: (1) antisense compounds, CC mPGES-1, which specifically hybridise with the nucleic acid encoding CC mPGES-1, which specifically hybridise with the nucleic acid encoding CC mPGES-1 in cells or tissues; and (3) a method of inhibiting the expression of CC mPGES-1 in cells or condition associated with mPGES-1. MPGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, antisinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, CC antisinflammatory, neuroprotective, nootropic, antistributies, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's CC condition associated with mPGES-1 ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                           Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-SEP-2002; 2002US-0413549P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human oligonucleotide #1846.
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                                                                                                            11-MAR-2004
                                                                                                                                                                             US2004049022-A1
                                                                                                                                                                                                                                             Homo sapiens
                                       25-JUL-2003; 2003US-00627930.
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AC ADO4
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T15-1
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DE Huma
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KW CCR1
KW CCR1

15-JUL-2004

(first entry)

ADO45362;

Human oligonucleotide

**#728.** 

Human, ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase & tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease;

RESULT 1126

ADO45362 standard; DNA;

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                                                                                                                                                                                                                                       The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region codon, coding recipion, 5' or 3' intron-exon junction, intron or region codon, coding recipion, 5' or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-6 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention code is relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the codigonucleotides are useful for reducing or inhibiting expression of a congene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, cCCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are cuseful for preventing or treating a respiratory or lung disease. The cc respiratory or lung disease is associated with hyper-responsiveness to inflammation or an inflammatory disease. The respiratory or lung disease is consen from airway inflammatory disease. The respiratory or lung disease (CCP), chronic obstructive pulmonary disease (COPD), calleric thinite archive respiratory distress sundrome. pollognary
                                                                    Matches
                                                                                   Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
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(SAND/)
(TANG/)
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23-APR-2002; 2002WO-US013143
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Shahabuddin S, Lu H, Cong
                                                                                                                                                                                           allergic rhinitis, acute respiratory distress syndrome, pulmonan hypertension, lung inflammation, bronchitis, airway obstruction bronchoconstriction. This sequence represents an oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 2; SEQ ID NO 1847; 174pp; English.
                                                                                                                                       Sequence 20
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                        352 CTCCTGAGCTCAAGCAGTC 370
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                                                                  l Similarity 94.
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MILLER S.
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LU H.
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 CTCCTGAGCTTAAGCAGTC 20
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                                                                                                                                         6 C; 4 G;
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ong H;
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Matches 18
                                                                                                                                                                                                                                                                                                                                                                                    gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, Ebtaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C, PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The respiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A receptor(s), and/or asthma and/or lung allergies associated with inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary disease (COPD),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL)-5 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 B, PDE4 B, PDE4 C or PDE4 D. The invention also relates to a method of screnning a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The oligonucleotides are useful for reducing or inhibiting expression of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   asthma; lung allergy; inflammation; inflammatory disease; airway inflammatton; allergy; impeded respiration; cystic fibro chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension;
                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; SEQ ID NO 728; 174pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-293804/27.
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23-APR-2002; 2002WO-US013143.
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                                                                                                                                                                                                                                                                                                     allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction obsoronchoconstriction. This sequence represents an oligonucleotide of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TANG/)
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AGUILAR D.
MILLER S.
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                                                                                                                                   Similarity
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                                                 GAGTAGCTGGGACTACAGG 746
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                                                                                                                                                                                                                      BP;
                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hyper-responsiveness; adenosine; adenosine A
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                                                                                                                                                                                                                         Α,
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Cong
                                                                                                                              1.8%;
                                                                                                                                                                                                                           ω
                                                                                                                                                                                                                         C; 8
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ong H;
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                                                                                                                                                                                                                         G; 4 T; 0 U; 0 Other
                                                                                                                                   Score 17.4;
Pred. No. 1
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                                                                                                            Mismatches
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                                                                                                                                   1.5e+03;
                                                                                                                                                              DB 1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cystic fibrosis;
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                                                                                                         Gaps
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RESULT 1127
ADO45271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; CODD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
Novel single or multiple target oligonucleotide anti-sense to e.g. Conitiation codon, intron of respiratory disease-relevant gene e.g. Coranness, MCP4, useful for prophylaxis or treating respiratory disease
                                                                                                                                               Shahabuddin S, Lu H,
                                                                                                                                                                     Nyce JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase B; PDE4 B; PDE4 C; PDE4 D; respiratory disease; tryptase b; ppE4 A; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AD045271;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADO45271 standard; DNA;
                                                                                                                                                                                                                                                                                                                                        (SAND/)
(TANG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-JUL-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-JUL-2004
                                                                                                                                                                                                                      (CONG/)
                                                                                                                                                                                                                                                                                                                                                                                         (NYCE/)
                                                                                                                                                                                                                                                                                           (MILL/)
                                                                                                                                                                                                                                                                     SHAH/)
                                                                                                                                                                                                                                                                                                                    AGUI/
                                                                                                                                                                                                                                                                                                                                                                                         NYCE
                                                                                                                                                                                                                      CONG H.
                                                                                                                                                                                                                                          H DT
                                                                                                                                                                                                                                                                                         MILLER S.
                                                                                                                                                                                                                                                                                                                                          TANG L.
                                                                                                                                                                                                                                                                SHAHABUDDIN S
                                                                                                                                                                                                                                                                                                                                                              SANDRASAGRA A.
                                                                                                                                                                                                                                                                                                                    AGUILAR D.
                                                                                                                                                                       Sandrasagra A,
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2002WO-US013143.
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                                                                                                                                               Cong
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                                                                                                                                                 Tang L,
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                                                                                                                                                                       Aguilar
                                                                                                                                                                          Á
                                                                                                                                                                          Miller
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CCR1, e.g.

Claim 2; SEQ ID NO 637; 174pp; English.

The invention relates to oligonucleotides anti-sense to an initiation CC codon, coding region, 5' or 3' intron-exon junction, intron or region CC with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target CC chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-6 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 CD 7 DE4 D. The invention CC also relates to a method of screening a candidate compound that binds to CC one or more nucleic acid target(s) or expressed product(s), for the CC prevention and/or treatment of a respiratory or lung disease. The CC constant of the constant of a respiratory or lung disease a, CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The CC crepiratory or lung disease is associated with hyper-responsiveness to CC and/or increased levels of, adenosine and/or levels of adenosine A creceptor(s), and/or asthma and/or lung allergies associated with cC inflammation or an inflammatory disease. The respiratory or lung disease

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RESULT 1128
AD052269/c
ID AD05226
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                             cytostatic; gene therapy; inhibitors of apoptosis-like; IAP-like; IAP-like modulator; IAP-like associated disorder; hyperproliferative disorder; human; antisense oligonucleotide; antisense technology; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction obronchoconstriction. This sequence represents an oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cystic fibrosis (CF), allergic rhinitis, acu
                                                                                                                                                                                                                                                                                                                                                       modified
                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human inhibitor of apoptosis-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADO52269
                                                                        New compound targeted to a nucleic acid molecule encoding inhibitors apoptosis (IAP)-like and inhibits expression of IAP-like, useful for modulating the expression of IAP-like or for treating, e.g.
                                                                                                                                                                                                                                27-MAY-2004.
                                                                                                                                                                                                                                                                                                            modified_base
 where the compound
                                                                                                                                                                                     22-NOV-2002;
                                                                                                                                                                                                          22-NOV-2002; 2002US-00303325
                                                                                                                                                                                                                                                      US2004102395-A1
                     The invention
                                          Example 14; SEQ ID NO 143; 58pp; English
                                                                hyperproliferative
                                                                                                                      WPI; 2004-399725/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    214 GTCTCGAACTCCCGACCTC
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                                                                                                                                                               ISIS
                                                                                                                                          CF,
                                                                                                                                                                                                                                                                                                                                                     _base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GTCTCGAACTCCTGACCTC
            acid molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                         Dobie KW;
                                                                                                                                                                                      2002US-00303325
                                                                                                                                                                PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
describes a compound 8-80 nucleobases in length targeted d molecule encoding inhibitors of apoptosis (IAP)-like, pound specifically hybridises with the nucleic acid molecu
                                                                                                                                                                                                                                                                                               /note= "
15. .20
/*tag=
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                                                                                                                                                                                                                                                                             note=
                                                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "OTHER= Phosphorothioate
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                  mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                 disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8 C;
                                                                                                                                                                                                                                                                                       base=
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ი
                                                                                                                                                                                                                                                                            "OTHER= 2'-O-Methoxyethyl
                                                                                                                                                                                                                                                                                                                       "OTHER=
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HER= 2'-O-Methoxyethyl
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Pred. No. 1.
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U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 Other;
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                                                                                                                                                                                                                                                                                                                                                                              backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                             (2'-MOE)
                                                                                                                                                                                                                                                                                                                        (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                                             nucleotides'
                                                                                                                                                                                                                                                                                                                        nucleotides"
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                                                                                                                                                                                                                                                                                                                                                                              cytidines
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le of the
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molecule

New compound targeted to apoptosis (IAP)-like and

a nucleic acid molecule encoding inhibits expression of IAP-like,

inhibitors useful for

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RESULT 1129
ADO52207
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Best Local s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            encoding IAP-like comprising 16000 bp (SEQ ID NO. 4) and inhibits the expression of IAP-like. Also described are: inhibiting the expression of IAP-like; and cells or tissues; screening for a modulator of IAP-like; a diagnostic method for identifying a disease state comprising identifying the presence of IAP-like in a sample using at least one of the primers selected from 2 sequences comprising SEQ ID NO. 5 or 6, or the probe comprising SEQ ID NO. 7, a kit or assay device comprising the compound; and treating an animal having a disease or condition associated with IAP-like. The compound is useful for modulating the expression of IAP-like. It is also useful for diagnosing or treating diseases associated with expression of IAP-like, e.g. a hyperproliferative disorder. This sequence represents a human inhibitor of apoptosis (IAP)-like antisense
                                                                                                                                                                                                                                                                                                                                                                                                    cyrostatic; gene therapy; inhibitors of apoptosis-like; IAP-like modulator; IAP-like associated disorder; hyperproliferative disorder; human; antisense olicomicla antisense technology: ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 3 A; 7 C; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human inhibitor of apoptosis-like antisense oligonucleotide seqid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AD052207
                                                                                                     22-NOV-2002;
                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                                                                                                                          Key
                                                                                                                                                                                                                                                                                                                                                                                   Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADO52207;
                                                          Bennett
                                                                                                                              22-NOV-2002;
                                                                                                                                                     27-MAY-2004
                                                                                                                                                                             US2004102395-A1
                                                                                (ISIS-)
                                                                                                                                                                                                                                                                                                                                                                                  sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                390 AAGTGCTGGGATTACAGGC 408
                                   2004-399725/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
18; Conserv
                                                          CF,
                                                                                ISIS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAGTGCTGGGATCACAGGC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                 PHARM INC
                                                                                                        2002US-00303325
                                                                                                                               2002US-00303325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                        Dobie KW
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/mod_base= OTHER
/mod_base= 2/
                                                                                                                                                                                                                                                                                                 are
                                                                                                                                                                                                                        /*tag=
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                                                                                                                                                                                                                                                                                                            note= "OTHER= Phosphorothicate backbone.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.8%;
94.7%;
                                                                                                                                                                                                                                                                                                 5-methylcytidines"
                                                                                                                                                                                                      "OTHER=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20
                                                                                                                                                                                            OTHER
HER= 2'-O-Methoxyethyl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 17.4;
Pred. No. 1
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); Mismatches
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                                                                                                                                                                                                                                                     -O-Methoxyethyl (2'-MOE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                      oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                       (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                              IAP-like;
                                                                                                                                                                                                                                                                                                               All cytidines
                                                                                                                                                                                                       nucleotides"
                                                                                                                                                                                                                                                      nucleotides"
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AD052271/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention describes a compound 8-80 nucleobases in length targeted to a nucleic acid molecule encoding inhibitors of apoptosis (IAP)-like, where the compound specifically hybridises with the nucleic acid molecule encoding IAP-like comprising 16000 bp (SEQ ID NO. 4) and inhibits the expression of IAP-like. Also described are: inhibiting the expression of IAP-like in cells or tissues; screening for a modulator of IAP-like; a cidagnostic method for identifying a disease state comprising identifying the presence of IAP-like in a sample using at least one of the primers selected from 2 sequences comprising SEQ ID NO. 5 or 6, or the probe comprising SEQ ID NO. 7; a kit or assay device comprising the compound; and treating an animal having a disease or condition associated with IAP-like. The compound is useful for modulating the expression of IAP-like. It is also useful for diagnosing or treating diseases associated with expression of IAP-like, e.g. a hyperproliferative disorder. This sequence represents a human inhibitor of apoptosis (IAP)-like antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modulating the expression of IAP-like or for treating, hyperproliferative disorder.
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                                                                                                                                                                                                                                                                                                                                                                                           cytostatic; gene therapy; inhibitors of apoptosis-like; IAP-like; IAP-like modulator; IAP-like associated disorder; hyperproliferative disorder; human; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human inhibitor of apoptosis-like antisense oligonucleotide seqid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-AUG-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 oligonucleotide.
             22-NOV-2002; 2002US-00303325
                                         22-NOV-2002; 2002US-00303325
                                                                                                  US2004102395-A1
                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                               antisense technology;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     μ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGCTCAAGCGATTCTCCTG 1015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                     /*tag=
                                                                                                                                                                          /note=
15. .20
                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                are 5-methylcytidines"
                                                                                                                                                                                                                                                           /mod_base= OTHER
/note= "OTHER= Phosphorothioate backbone. All cytidines
                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                  *tag=
                                                                                                                                                                                                    mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ŏ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.8%;
                                                                                                                                                                          . 20
                                                                                                                                             base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 81; 58pp;
                                                                                                                                                                                      "OTHER=
                                                                                                                               "OTHER=
                                                                                                                                                            ი
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 17.4;
Pred. No. 1.
                                                                                                                            2'-0-Methoxyethyl (2'-MOE)
                                                                                                                                                                                      2'-O-Methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 20;
                                                                                                                                                                                      (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                               nucleotides"
                                                                                                                                                                                      nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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RESULT 1131
ADO52203
ID ADO5220
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Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diagnostic method for identifying a disease state comprising identifying the presence of IAP-like in a sample using at least one of the primers selected from 2 sequences comprising SEQ ID NO. 5 or 6, or the probe comprising SEQ ID NO. 7, a kit or assay device comprising the compound; and treating an animal having a disease or condition associated with IAP-like. The compound is useful for modulating the expression of IAP-like. It is also useful for diagnosing or treating diseases associated with expression of IAP-like, e.g. a hyperproliferative disorder. This sequence represents a human inhibitor of apoptosis (IAP)-like antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New compound targeted to a nucleic acid molecule encoding inhibitors apoptosis (IAP)-like and inhibits expression of IAP-like, useful for modulating the expression of IAP-like or for treating, e.g. hyperproliferative disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention describes a compound 8-80 nucleobases in length targeted to a nucleic acid molecule encoding inhibitors of apoptosis (IAP)-like, where the compound specifically hybridises with the nucleic acid molecule encoding IAP-like comprising 16000 bp (SEQ ID NO. 4) and inhibits the expression of IAP-like. Also described are: inhibiting the expression of IAP-like, also described are: inhibiting the expression of IAP-like; a screening for a modulator of IAP-like; a
                                                                                                                                                                                                                                                                                cytostatic; gene therapy; inhibitors of apoptosis-like; IAP-like; IAP-like modulator; IAP-like associated disorder; hyperproliferative disorder; human; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                           Human inhibitor of apoptosis-like antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 14; SEQ ID NO 145; 58pp; English.
                                                                                                                                                                                                           ey
                                                                                                                  modified_base
                                                                                                                                                                                                                                         Homo
                                                                                                                                                                                                                                                                     antisense technology; ss.
                                                                                                                                                                                                                                                                                                                                                                          12-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                          ADO52203;
                                                                                                                                                                                                                                                                                                                                                                                                                                      ADO52203 standard;
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                                                      modified_base
                                                                                                                                                                                           modified_base
                                                                                                                                                                                                                                       sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGTTCAAGCGATTCTCCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGCTCAAGCGATTCTCCTG 1015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                      /note= 15. .20
                                                                                                                                 are
                          /*tag=
/mod_ba
                                                                                                                                                 /mod_base= OTHER
/note= "OTHER= P
                                                                                                                                                                                                         Location/Qualifiers
                                                                                       /mod
                                                                                                  /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                5-methylcytidines"
                                                                      _base= OTHER
e= "OTHER= 2'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 C; 6 G; 3 T;
                            base= OTHER
               "OTHER=
                                                                                                                                                                                                                                                                                                                                                                                                                                       20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 17.4;
Pred. No. 1
              2'-O-Methoxyethyl (2'-MOE) nucleotides'
                                                                                                                                                  Phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                       -O-Methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                              oligonucleotide seqid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                         (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                         nucleotides'
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                                                                                                                                                  cytidines
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RESULT 1132
ADP45826
ID ADP4582
XX Brideric Advance and Advance an
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New compound targeted to a nucleic acid molecule encoding inhibitors of apoptosis (IAP)-like and inhibits expression of IAP-like, useful for modulating the expression of IAP-like or for treating, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-MAY-2004
                                                                                                                                                                                                                                                                             breast cancer; cytostatic; gene therapy; human; intercellular adhesion molecule, ICAM-1; human; receptor; BB2; CD54; cell surface glycoprotein P3.58; ICAM-4; Landsteiner-Wiener blood group; ICAM-5; telencephalin; chromosome 19p13; ss; primer; PCR; SNP; single nucleotide polymorphism; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-NOV-2002; 2002US-00303325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-NOV-2002; 2002US-00303325.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2004102395-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADP45826;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADP45826 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 14; SEQ ID NO 77; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hyperproliferative disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bennett
                                      25-NOV-2003; 2003WO-US037948.
                                                                                                    10-JUN-2004.
                                                                                                                                                           WO2004047623-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2004-399725/37
                                                                                                                                                                                                                         sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAGTGCTGGGATTACAGGC 408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAGTGCTGGGATCACAGGC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18 used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 C; 7 G; 3 T; 0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   to genotype human ICAM-1/ICAM-4/ICAM-5 SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 17.4; DB 1
Pred. No. 1.5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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of breast cancer comprising detecting the presence of absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a subject at risk of breast cancer, for early diagnosis, prevention and treatment of breast cancer, possibly via gene therapy, as well as to analyse and predict a response to a breast cancer treatment and in clinical druy trials. The current sequence is that of an Extend primer (also described as probe) of the invention which was used to genotype human intercellular adhesion molecule ICAM-1/ICAM-4/ICAM-5 gDNA. ICAM-1 (human rhinovirus receptor; BB2; CD54; cell surface glycoprotein P3.58) has been mapped to chromosomal position 19p13.3-p13.2, ICAM-4 (Landsteiner-Wiener blood group; LW) has been mapped to chromosomal position 19p13.3-pha been mapped to chromosomal position 19p13.3-pha been mapped to chromosomal position 19p13.2-cen and ICAM-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying a subject at risk of breast cancer by detecting the presence of polymorphic variations in the ICAM, MAPK10, KIAA0861, NUMA1 or GALE regions which are associated with breast cancer in a nucleic acid sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-NOV-2002; 2002US-0429136P.
24-JUL-2003; 2003US-0490234P.
Sequence 20
                                                                                                                                                                                                                                                                                                                                                                 The
                                                                                                                                                                                                                                                                                                                                                                                                         Example 4; Page 83; 289pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-441051/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Roth RB
                                                                                                                                                                                                                                                                                                                                                                                                                                                     from a subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (SEQU-)
                                            (telencephalin)
                                                                                                                                                                                                                                                                                                                                                                 invention relates to a novel method for identifying a subject at risk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENOM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nelson MR,
BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                            has been mapped to chromosomal position 19p13.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Braun
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Matches
                                     Query Match
Best Local (
               635
                               18;
N
                                      Similarity
           CTCTGTCACCCAGGCTGGA 653
CTTTGTCACCCAGGCTGGA
                               Conservative
                                     1.8%;
                               0
20
                                       Pred.
                                              Score 17.4;
                                Mismatches
                                        No.
                                       1.5e+03;
                                               DB 1;
                                              Length
                                Indels
                                                20;
                                <u>,</u>
                                Gaps
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RESULT 1133
AAQ10789
                                                              09-AUG-1989;
                                                                                                        Synthetic
                                                                                                                     transgenic
                                                                                                                           Human factor IX; blood clotting; trans-immortalised cell lines;
                                                                                                                                         Probe for identifying cDNA clones encoding human factor IX.
                                                                                                                                                         25-MAR-2003
08-MAY-1991
                                                                                                                                                                               AAQ10789;
                                                                                                                                                                                             AAQ10789 standard;
                                                 09-AUG-1989;
                                                                            21-FEB-1991.
                                                                                          WO9102056-A
                                  (TRGE ) TRANSGENE
                     1991-073532/10.
                                                                                                                      animals; type
                                                                                                                                                         (revised)
(first entry)
                                                89FR-00010720
                                                               89FR-00010720
                                   SA
                                                                                                                                                                                             DNA;
                                                                                                                                                                                              21
                                                                                                                      W
                                                                                                                     haemophilia;
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are New

immortalised cell lines expressing biologically active factor-IXsi obtd. from new transgenic(s) with human factor-IX-expressing DNA

fragment incorporated into their

1; Page 8;

37pp;

French

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Best Local S
Matches 18
Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This 21 mer probe is used to screen a human lymphoblastoid cell line-derived lambda EMBL3 genomic library. The positive clones obtd. are sequenced and their overlapping sequence information is used to prepare synthetic DNA sequence used in the prepn. of recombinant human factor IX A trans-immortalised cell line with the ability to express human factor IX can be produced as can transgenic animals having this exogenous DNA fragment integrated into their genomes. The recombinant human factor IX is useful in the treatment of type B haemophilia. See also Q10784-88 and Q10853-63. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH37857 standard;
                                                                                                                                                                                                                            New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP specific upper PCR primer SEQ ID
                                                                                                                                                                          Claim 1; Page 53; 83pp; English
                                                                                                                                                                                                                                                                                                              Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                 15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200129262-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Single nucleotide polymorphism; SNP; single nucleotide primer extension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                   13-OCT-2000; 2000WO-US028436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sapiens
                                                                                                                                                                                                                                                                                2001-290930/30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ب
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                                                                                                                                                                                                                                                                                                                                                ORCHID BIOSCIENCES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GATTACAGGCGTGAGCCAC 887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GATTATAGGCGTGAGCCAC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 5 A; 4 C; 7 G; 5 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   forensic investigation; paternity analysis; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                 99US-0160096P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             뫄
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           653.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1:
                                                                                                                                                                                                                            polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IX.
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RESULT 1135
AAH38405/c
ID AAH3840
XX AAH3840
XX SAH3840
XX Single
XX Lesch.N
XX Lesch.N
XX Inflamm
XX Homo 88
XX WO20012
XX WO20012
XX WO20012
XX For 13-OCT.
XX I5-OCT.
XX IF-OCT.
XX PP 13-OCT.
XX PP 13-OCT.
XX PP 13-OCT.
XX PF 13-OCT.
XX PF 13-OCT.
XX Single
PM WO20012
XX WO20012
XX Single
PM WO20012
XX CORCH-)
XX PF 13-OCT.
XX Single
PM A CORCH-)
XX ORCH-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a PCR primer specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 6 A; 6 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                        New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Single nucleotide polymorphism; SNP; single nucleotide primer extension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP specific upper PCR primer SEQ ID 1201.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH38405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-APR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      identity of a SNP and for genotyping nucleic acid samples, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tor a human SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                               2001-290930/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORCHID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GTCAGGCTGGTCTCGAACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000WO-US028436.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 forensic investigation; paternity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-0160096P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    containing DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pohl M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        223
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 17.4; DB 1;
Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ٥.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by

a single-nucleotide

primer extension

Sequences AAH37205 - AAH40944 represent PCR primers,

single nucleotide

Claim 1; Page 56; 83pp; English.

sample.

В S

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RESULT 1136
AAF24290
ID AAF2429
                                                                      Query Match
Best Local S
Matches 18
Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syddrome, muscula dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diseases, including, rheumatold architatis, muture everyone inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and microorganism. The method is also useful in forensic investigations and microorganism. The method is also useful in forensic investigations and microorganism.
                                                                                                                                                                                                                                          Determining complementarity of nucleotide fragment for gene analysis, comparing flow of electric current from or to electroconductive subst through DNA fragment, with reference obtained from its complement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Complementary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF24290
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 5 A; 6 C; 4 G; 5 T; 0 U; 1 Other;
                                                                                                                    The present invention provides a method for analysing a nucleic acid strand to determine the degree of complementarity between two sequences. This involves the measurement of an electric current along the annealed strands compared to a standard. This is useful in the analysis of genetic strands compared to a standard. This is useful in the analysis of genetic strands compared to a standard.
                                                                                                                                                                                                           Example 1; Page 12; 28pp; English.
                                                                                                                                                                                                                                                                                                                    WPI; 2001-140003/15.
                                                                                                                                                                                                                                                                                                                                                                                                                            07-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-JUN-2000; 2000EP-00112235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EP1065278-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Complementary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF24290
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          paternity analysis. The present sequence represents a
                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        chip;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1086 AGAGGCGGGGTTTCACCATAT 1106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGAGAYGGGGTTTCACCATCT 1
                                                                      21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer; ss.
                                                                                                                                                                                                                                                                                                                                                     Abe Y,
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                                                                      BP; 1
   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic
                                                                                                       and variation
                                                                                                                                                                                                                                                                                                                                                                                                                          99JP-00159339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      containing DNA sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                     Ogawa M,
                1.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          acid; gene analysis; polymorphism; variation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             acid
                                                                      c; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             detection method related
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                                                                                                       between
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                                                                                                                                                                                                                                                                                                                                                     Takagi M,
 Score 17.4; DB 1;
Pred. No. 1.5e+03;
0; Mismatches 1
                                                                      20 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                         Takenaka
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                                                                      0 Other;
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                                   Length
   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence
                                                                                                                                                                                                                                                                                                                                                         Yamashita
                                                                                                                        analysis of genetic
                                    21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for e.g. to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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                                                                                                                                                                                                                                                                lysis, by substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            specific
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RESULT 1138
ABS60598/c
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                                                                                                                                                                                                                                                                                                                                                                  RESULT 1137
ABK88537/c
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                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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         Aminopeptidase P; XPNEP2; bradykinin receptor B1; ds; BDKRB1; tachykinin receptor B1; TACR1; C1 esterase inhibitor; C1NH; kallikrei: KLK1; bradykinin receptor B2; BDKRB2; gene therapy; angiotensin converting enzyme 2; ACB2; protease inhibitor 4; PI4; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; angioedaema; aneurysm; embolism; thrombosis; coronary artery disease; angioedaema;
                                                                                                                                                                                                                                                                                                                                                                                                    The invention describes a method of diagnosing a panic disorder with a polymorphism of the upper stream region of human cholecystokinin (CCK) gene. This sequence represents a human cholecystokinin gene associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Diagnosis and identification of panic disorder caused upper stream region of human cholecystokinin gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human cholecystokinin associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK88537;
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                                                                                                                                                                                                                                                                                                                                                                 Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-569888/61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-DEC-2000; 2000JP-00375090
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Panic disorder;
arteriosclerosis;
                                                                                                                       Human polymorphism associated DNA sequence #347.
                                                                                                                                                 05-NOV-2002
                                                                                                                                                                         ABS60598;
                                                                                                                                                                                                 ABS60598 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 6; Page 6; 13pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RIKA ) RIKAGAKU KENKYUSHO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-JUN-2002.
                                                                                                                                                                                                                                                                                                                                                                                          PCR primer
                                                                                                                                                                                                                                                                                                                                          Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       427
                                                                                                                                                                                                                                                                                       645
                                                                                                                                                                                                                                                                21
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                                                                                                                                                                                                                                                                CAGGCTGGAGTACAGTGGC 3
                                                                                                                                                                                                                                                                             CAGGCTGGAGTGCAGTGGC 663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TTTTTTTTTATTTTTTT
                                                                                                                                                                                                                                                                                                                                                                 BP; 4 A; 8 C; 4 G; 5 T;
                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                 (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
                                                                                                                                                 entry)
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                                                                                                                                                                                                                                                                                                                 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         445
                                                                                                                                                                                                                                                                                                                                         Score 17.4;
                                                                                                                                                                                                                                                                                                                             Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human cholecystokinin; upper stream; CCK;
                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer P1
                                                                                                                                                                                                                                                                                                                                                                   0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                                                                                            21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polymorphism
                                                                                                                                                                                                                                                                                                                 0;
                                                                                       kallikrein
                                                   trachoma;
                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of.
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CC tachykinin receptor B1 (RARRI), C1 esterase inhibitor (CHH), kallikrein C2 (ACE2) or protease inhibitor 4 (PIA), comprising at least one C2 (ACE2) or protease inhibitor 4 (PIA), comprising at least one C3 (ACE2) or protease inhibitor 4 (PIA), comprising at least one C4 (PIA), comprising at least one C5 (PIA) probe that hybridies to a C5 (PIA) position as provided in the detailed summary of single C6 (PIA) probe that hybridies to a C6 (PIA) probe that hybridies to a C7 (PIA) probe that hybridies to a C7 (PIA) probe that hybridies to a C8 (PIA) probe that hybridies to a C9 (PIA) probe that hybridies to a C9 (PIA) probe the sample comprising probe the sample comprising probe the sample comprising probe that in a probe the polymorphic positions in a gene encoding a protein selected from the group above; (3) constructing (M2) C8 (PIA) identifying (M3) an individual at risk of developing a disorder C9 (PIA) identifying (M3) an individual at risk of developing a disorder C9 (PIA) protein selected from the group above; and (6) genotyping (M4) and C9 (PIA) protein selected from the group above; and (6) genotyping (M4) and C9 (PIA) protein selected from the group above; and (6) genotyping (M4) and C9 (PIA) protein are useful for detecting, diagnosing, treating the C9 (PIA) protein are useful for detecting, diagnosing, treating at C9 (PIA) protein protein selected such as angloedaem and diseases which C9 (PIA) protein protein selected such as angloedaem and diseases which C9 (PIA) protein protein selected such as angloedaem and diseases which C9 (PIA) protein selection, ventricular hypertrophy, vascular diseases, aneutysm, embolism, thrombosis, coronary C1 (PIA) protein selection protein 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tsuchihashi
Swanson BN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-DEC-2000; 2000US-0251015P.
23-JAN-2001; 2001US-0263678P.
02-MAR-2001; 2001US-0273037P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD; Chronic obstructive pulmonary disease; enterocolitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 812;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-619265/66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-DEC-2001; 2001WO-US047235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200261131-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to an isolated nucleic acid from a human gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (BRIM )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          isolated nucleic acid with at least one polymorphic position, useful detecting, diagnosing and treating disorders such as angioedema, cer, viral, bacterial or fungal infection, cardiovascular and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ) BRISTOL-MYERS :
) TSUCHIHASHI Z.
) HUI L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              aminopeptidase P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             diseases.
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Powell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        977pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bradykinin receptor B1
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Best Local Similarity

1.8%; 2 C; 7

Score 17.4; DB 1; Pred. No. 1.5e+03;

Length 21

Query Match Sequence 21

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RESULT 1139
ABS60817/c
ID ABS6081
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            i (KIKI), bradykinin receptor B2 (BDKRB2), angiotensin converting enzyme 2 (ACB2) or protease inhibitor 4 (PI4), comprising at least one polymorphic position. Also included are (1) a probe that hybridises to a polymorphic position as provided in the detailed summary of single nucleotide polymorphisms comprising additional 5 and 3 flanking genomic sequence; (2) analysing (M1) at least one nucleic acid sample comprising obtaining the sample from one or more individuals and determining the nucleic acid sequence at one or more polymorphic positions in a gene encoding a protein selected from the group above; (3) constructing (M2) haplotypes using the genes comprising grouping at least two nucleic acids; (4) identifying (M3) an individual at risk of developing a disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; aneurysm; embolism; thrombosis; corronary artery disease; angioedaema; arteriosclerosis; atherosclerosis; hypersensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tachykinin receptor B1; TACR1; C1 esterase inhibitor; C1NH; 
KKK1, bradykinin receptor B2; BDKBB2; gene therapy;
angiotensin converting enzyme 2; ACE2; protease inhibitor 4;
polymorphism; haemangioma; tumour; sarcoma; Crohn's disease;
                                                                                                                                                                                                          The invention relates to an isolated nucleic acid from a human gene encoding aminopeptidase P (XPNEP2), bradykinin receptor B1 (BDKRB1) tachykinin receptor B1 (TACR1), C1 esterase inhibitor (C1NH), kalli
                                                                                                                                                                                                                                                                                                                                     New isolated nucleic acid with at least one polymorphic position, useful for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and
                                                                                                                                                                                                                                                                                                                                                                                                                                           Tsuchihashi Z, Hui L, Swanson BN, Powell JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TSUC/)
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23-JAN-2001;
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                                                                                                                                                                                                                                                                                  Disclosure; Page 884; 977pp; English
                                                                                                                                                                                                                                                                                                                      autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-619265/66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BRIM )
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) identifying (M3) an individual at risk of developing a administration of an ACE inhibitor and/or vasopeptidase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TSUCHIHASHI Z.
HUI L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BRISTOL-MYERS SQUIBB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                obstructive pulmonary disease;
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                                                                                                                                                                                                                                                                                                                      diseases
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2001US-0263678P.
2001US-0273037P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ma-Edmonds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protease inhibitor 4; PI4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         trachoma;
                                                                                                                                                                                                                kallikrein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC using the polymorphic data; (5) a library of nucleic acids, each of which CC comprises one or more polymorphic positions within a gene encoding a CC human protein selected from the group above; and (6) genotyping (M4) an CC individual comprising obtaining a nucleic acid sample, determining the CC nucleotide present in at least one polymorphic position, and comparing at CC least one position with a known data set. The genes, (M1, M2, M3 and M4) CC and compositions are useful for detecting, diagnosing, treating, CC preventing various disorders such as angioedaema and diseases which CC involve angiogenesis like haemangiomas, tumours, sarcomas, Crohn's CC disease, trachomas, and cardiovascular diseases like angina pectoris, CC hypertension, heart failure, myocardial infarction, ventricular CC hypertension, heart failure, myocardial infarction, ventricular CC hypertension, heart failure, myocardial infarction, ventricular CC hypersensitivity reactions, sepsis, autoimmune diseases, inflammatory CC arthritis, cancer, wounds, viral, bacterial or fungal infection, Chronic Obstructive pulmonary disease (COPD) and enterocolitis (many other CC diseases and disorders are listed in the specification). The CC against the proteins may be utilised for immunophenotyping of cell lines and biological samples. The present sequence is included in the sequence of listing but is not referred to anywhere plac in the securification.
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ABS60599/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1140
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Best Local S
Matches 18
                                                                                                            04-DEC-2000;
23-JAN-2001;
02-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                      angiotensin converting enzyme 2; ACE2; protease inhibitor 4; PI4; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; aneurysm; embolism; thrombosis; coronary artery disease; angioedaema; arteriosclerosis; atherosclerosis; hypersensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Aminopeptidase P; XPNEP2; bradykinin receptor B1; ds; BDKRB1; tachykinin receptor B1; TACR1; C1 esterase inhibitor; C1NH; kallikrein KLK1; bradykinin receptor B2; BDKRB2; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABS60599 standard; DNA; 21
                                                                                                                                                                                      03-DEC-2001; 2001WO-US047235.
                                                                                                                                                                                                                           08-AUG-2002.
                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                            Chronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human polymorphism associated DNA sequence #348.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1056 CCACACCCCGCTAATTTTT 1074
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19 CCACACCCAGCTAATTTTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                         BRISTOL-MYERS
TSUCHIHASHI Z.
                                                                                                                                                                                                                                                                                                                                            obstructive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                            2000US-0251015P.
2001US-0263678P.
2001US-0273037P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 samples. The present sequence is included in the sunot referred to anywhere else in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.8%;
94.7%;
                                                                                                                                                                                                                                                                                                                                        pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N
                                                                           SQUIBB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ç;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 17.4;
Pred. No. 1.
                                                                             8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.5e+03;
                                                                                                                                                                                                                                                                                                                                            enterocolitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   trachoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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Tsuchihashi Z,

Hui L,

Zerba

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Ma-Edmonds M,

Perrone

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TXAXEXEXEXEXEXE

05-NOV-2002

(first

entry)

ABS60816 standard; DNA;

21

ВP

Human polymorphism associated DNA sequence #453

Aminopeptidase P; XPNEP2; bradykinin receptor B1; ds; BDKRR1; tachykinin receptor B1; TACR1; C1 esterase inhibitor; C1NH; kallikrein KLK1; bradykinin receptor B2; BDKRB2; gene therapy; ALK1, bradykinin receptor B2; BCRB2; grotease inhibitor 4; PI4; angiotensin converting enzyme 2; ACB2; protease inhibitor 4; PI4;

밁 Ś

19

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Matches Query Match Best Local (

18;

Conservative

0; Pred.

Mismatches

0

Gaps

0

No. 1.5e+03;

Similarity

1.8%;

Score 17.4;

DB 1;

Length Indels

1056 CCACACCCCGCTAATTTT 1074

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CC (ACE2) or protease inhibitor 4 (P14), comprising at least one polymorphic position. Also included are (1) a probe that hybridises to a CC polymorphic position. Also included in the detailed summary of single converting enzyme (2) analysing (M1) at least one muclecide polymorphisms comprising additional 5 and 3 flanking genomic converting the sample from one or more individuals and determining the comprising obtaining the sample from one or more individuals and determining the converting from the group above; (3) constructing (M2) at least one nucleic acid sequence at one or more polymorphic positions in a gene concoding a protein selected from the group above; (3) constructing (M2) comprises one or more polymorphic positions at least two nucleic acids (2) (4) identifying (M3) an individual at risk of developing a discorder (2) upon administration of an ACE inhibitor and/or vasopeptidase inhibitor (2) (3) constructing (M2) and (3) identifying (M3) an individual at risk of developing a discorder (3) comprises one or more polymorphic positions within a gene encoding a comprise one or more polymorphic positions within a gene encoding a comprise one or more polymorphic positions within a gene encoding a comprise one position with a known data set. The genes, (M1, M2, M3 and M4) and compositions are useful for detecting, diagnosing, treating, (4) individual compositions are useful for detecting, diagnosing, treating, (4) individual various and comparing at composition with a known data set. The genes, (M1, M2, M3 and M4) and (4) compositions are useful for detecting, diagnosing, treating, (5) and compositions, and comparing at composition various such as angioedaema and diseases which (5) and composition with a known data set. The genes, (M1, M2, M3 and M4) and (5) composition with a known data set. The genes, (M1, M2, M3 and M4) and (5) composition are useful for detecting, diseases which (5) compositions are useful for detecting, composition, compositions, compositions, compositions, compositions, compositions, 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated nucleic acid with at least one polymorphic position, us for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             encoding aminopeptidase P (XPNE) tachykinin receptor B1 (TACR1),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated nucleic acid from a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 812; 977pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2002-619265/66
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            7
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        σ
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        Ή;
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        ۵,
            0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BDKRB1)
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Ct tachnykinin receptor Bi (YACKI), to escerable inhibitor (CINH), Kallikrein CC (CINH), Comprising at least one CC polymorphic position as provided in the detailed summary of single CC nucleotide polymorphisms comprising additional 5' and 3' flanking genomic sequence; (2) analysing (M1) at least one nucleic acid sample comprising CC obtaining the sample from one or more individuals and determining the nucleic acid sequence at one or more polymorphic positions in a gene CC encoding a protein selected from the group above; (3) constructing (M2) an individual at risk of developing a disorder CC using the polymorphic data; (5) a library of nucleic acids, each of which comprises one or more polymorphic positions within a gene encoding a nucleic acid sample, determining the nucleic data; (5) a library of nucleic acids, each of which CC comprises one or more polymorphic positions within a gene encoding a nucleic data; (5) a library of nucleic acids, each of which CC comprising obtaining a nucleic acid sample, determining the cC involve angiogenesis like haemangiomas, tumours, sarcomas, Crohn's CC involve angiogenesis like haemangiomas, tumours, sarcomas, Crohn's CC disease, trachomas, and cardiovascular diseases like angina pectoris, and compositions are useful for detecting, disgnosing, treating, pectoris, of hypertrophy, vascular diseases, aneuryme, embolism, thrombosis, coronary CC artery disease, arteriosclerosis and/or atherosclerosis, and compositions are also useful for chromsome infection. Antibodies coblymucleotides are also useful for chromsome identification. Antibodies colymucleotides are also useful for chromsome identification. Antibodies colymucleotides are also useful for chromsome identification.
against the proteins may be utilised for immunophenotyping of and biological samples. The present sequence is included in the listing but is not referred to anywhere else in the specificat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    encoding aminopeptidase P (XPNEP2), bradykinin receptor B1 (Bitachykinin receptor B1 (TACR1), C1 esterase inhibitor (C1NH),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     autoimmune diseases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-JAN-2001;
02-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-DEC-2001; 2001WO-US047235.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           aneurysm; embolism; thrombosis; coronary artery disease; angioedaema; arteriosclerosis; atherosclerosis; hypersensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cardiovascular disease; angina pectoris; hypertension; myocardial infarction; ventricular hypertrophy; vascula
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 884; 977pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention relates to an isolated nucleic acid from a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  isolated nucleic acid with at least one polymorphic position, useful detecting, diagnosing and treating disorders such as angioedema, cer, viral, bacterial or fungal infection, cardiovascular and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2002-619265/66
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      obstructive pulmonary disease; enterocolitis
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Powell JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000US-0251015P.
2001US-0263678P.
2001US-0273037P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Crohn's disease; trachoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        z
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   heart failure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     numan gene
(BDKRB1),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        kallikrein
                                                                                   cell lines
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Matches 18; Conserv
                                comprises detecting tandem repeats in a target coding sequence, scoring the repeats for polymorphic probability and generating a dataset correlating the repeats with polymorphic probability to identify a candidate polymorphic repeat. The computational methods (polymorphic marker prediction of ubiquitous simple sequences, POMPOUS, and Rep-X) are useful for identifying and detecting candidate polymorphic repeats in human genes, which can be used to understand, treat or eliminate genetic diseases, predispositions or adverse drug-treatment reactions. Examples of diseases linked to nucleotide repeats are Machado-Joseph, Haw River syndrome, Huntington's disease, fragile-X syndrome, Fredreich's ataxis, myotonic dystrophy, hyperandrogenaemia, spinal and bulbar atrophy and spinocerebellar ataxia. The sequences presented in ABX79676-ABX80022 are the polymorphic repeats identified for a search of human ESTs
                                                                                                                                                                                                                                                                                                                                                                Identifying a candidate polymorphic repeat within a coding sunderstanding or treating genetic disease, comprises detectivepeats in a target coding sequence and scoring the repeats
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EST; expressed sequence tag; ss; polymorphic repeat; tandem repeat; polymorphic marker prediction of ubiquitous simple sequences; POMPO Rep-X; human; genetic disease; drug-treatment; Machado-Joseph; Haw River syndrome; Huntington's disease; fragile-X syndrome; Fredreich's ataxis; myotonic dystrophy; hyperandrogenaemia; spinal atrophy; bulbar atrophy; spinocerebellar ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                           repeat within a coding sequence (expressed sequence tag, EST)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-APR-2003
                                                                                                                                                                                                                                                                                    The invention
                                                                                                                                                                                                                                                                                                                    Example; Col 495; 588pp; English.
                                                                                                                                                                                                                                                                                                                                                     polymorphic probability.
                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-208818/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Garner HR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EST polymorphic DNA repeat polynucleotide #119.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABX79794 standard; cDNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (TEXA ) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1056 CCACACCCCGCTAATTTT 1074
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 BP; 6 A; 2 C; 7 G; 6 T; 0 U; 0 Other.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CCACACCCAGCTAATTTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wren JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                  discloses a method for identifying a candidate polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-00475947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99US-00475947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                        polymorphic repeat within a coding sequence, 
ng genetic disease, comprises detecting tande
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 17.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fondon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     POMPOUS;
                                                                                                                                                                                                                                                                                                                                                                                            tandem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                              for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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Query Match
Best Local Similarity
Matches 18; Conserv

Conservative

0;

1.8%;

Score 17.4; DI Pred. No. 1.5e. 0; Mismatches

.5e+03 DB 1;

Length

Indels

٥.

Gaps

0

Sequence

21

₿P;

1 A; 0 C; 0

G; 20 T; 0 U; 0 Other;

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RESULT 1143
ADG79161
ID ADG7916
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                                                                                                                                                                       ABZ58551
ID ABZ
XX
AC ABZ
XX
AC ABZ
XX
DT 13-
XX
DE PCR
                                                                                                                 Ś
                                                                                                밁
                                                                      RESULT 1144
                                                                                                                                           Query Match
Best Local 9
                                                                                                                                     Matches
                                                                                                                                                                                       The invention comprises a method of diagnosing schizophrenia or a susceptibility to schizophrenia. The method involves detecting a polymorphism in a gene encoding a calcineurin (CN) subunit or CN-interacting molecule. The method of the invention is useful for the diagnosis of schizophrenia or a susceptibility to schizophrenia. The present DNA sequence represents a genotyping PCR primer that was used in an example of the invention.
                                                                                                                                                                                                                                                                                Diagnosing schizophrenia or susceptibility to schizophrenia comprises detecting a polymorphic variant of a polymorphism in a coding or non-coding portion of a gene encoding a calcineurin (CN) subunit or a CN interacting molecule.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               schizophrenia; polymorphism detection; calcineurin; CN; CN-interacting molecule; PCR; primer; ss; genotyping; PPP3CA; calcineurin A catalytic subunit-alpha.
                                                                                                                                                                                                                                                                                                                                                                                              26-MAR-2002; 2002US-0367944P.
07-MAR-2003; 2003US-0452813P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Calcineurin A catalytic subunit-alpha (PPP3CA) genotyping PCR primer #3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADG79161;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADG79161 standard;
                                                                                                                                                                       Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                          26-MAR-2003; 2003WO-US009578
                                                                                                                                                                                                                                                                                                                                                                                                                                            09-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2003082210-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified
                                                                                                                                                                                                                                                                 Example
PCR primer MR
                                                     ABZ58551
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          427
                                                                                                                  615
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                                                                                                N
                                                                                                                                                                                                                                                                                                                                                                   MASSACHUSETTS INST TECHNOLOGY.
UNIV ROCKEFELLER.
                                                                                                                                                                                                                                                                4; Page 173; 177pp; English.
                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TTTTTATTTTATTTTTTT 445
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TTTTTTTTTTTTTTTT
                                                     standard;
                                                                                                                  TTTTTGAGACAGAGTCTCA 633
                                                                                                TTTTGAGACGGAGTCTCA 20
                                                                                                                                                                       BP; 4 A; 4 C; 6 G; 7 T; 0
                                                                                                                                                                                                                                                                                                                                                  Karayiorgou M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                     Conservative
                   (first entry)
for diagnosis of Friedrich's ataxia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                     DNA;
                                                                                                                                           1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21
                                                      21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
                                                                                                                                                                                                                                                                                                                                                  Miyakawa T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20
                                                                                                                                   Pred. No. 1.56
); Mismatches
                                                                                                                                                     Score 17.4;
                                                                                                                                                                        ς,
                                                                                                                                             1.5e+03;
                                                                                                                                                                         0 Other;
                                                                                                                                                                                                                                                                                                                                                   Tonegawa
                                                                                                                                                     DB 1; Length 21;
                                                                                                                                      Indels
                                                                                                                                      0
                                                                                                                                      Gaps
                                                                                                                                      0
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RESULT 1145
ADPO8769/C
ID ADPO876
XX ADPO876
XX ADPO876
AC ADPO876
AC ADPO876
C Extend
XX Extend
XX Breast
KW Breast
KW GP6; GP
KW Single
XX Homo sa
XX Homo sa
XX WO20040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC multiplication disease of repeated trinucleotide sequence. The methods convolves amplification of the repeated trinucleotide sequence by PCR, cc analysis of the amplified product on microcapillary electrophoresis (CE), cc and determining the number of repeated trinucleotide repeats on the basis cc and determining the number of repeated trinucleotide repeats on the basis cc and determining the number of repeated trinucleotide is repeated 7-22 times in cc genetic region 9q13-q21.1, a GAA trinucleotide is repeated 7-22 times in cc genetic region 9q13-q21.1, a GAA trinucleotide is repeated 7-22 times in cc requence is that of reverse primer FR which is specific to the FA cc repeated trinucleotide sequence region. It is used with forward primer FR cc (see ABZ59550) to detect FA. A diagnosis kit comprising these primers is claimed. In a healthy subject, a PCR product of 157 bp is produced. Use cf CE, especially fabricated as an on-chip analysis system, allows the csize of the PCR product to be measured rapidly, with accuracy and cc reproducibility. The method allows diagnosis before the disease develops and determination of whether a silent carrier will develop the disease or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Diagnosing multiplication disease of repeated trinucleotide sequences e.g. Huntington's disease, by amplifying repeated trinucleotide sequence region, migrating and separating product by microcapillary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Friedrich's ataxia; diagnosis; microcapillary electrophoresis;
țrinucleotide repeat; screening; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-256603/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-AUG-2001; 2001KR-00047301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-AUG-2002; 2002WO-KR001489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2003014396-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a method for diagnosis of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 14; Page 8; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kim J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                               breast cancer; cytostatic; gene therapy; GP6; GPIV; GPVI; chromosome 19q13.4; ss; single nucleotide polymorphism.
                                                                                                                                                                                                     Extend primer 106 used to genotype human glycoprotein VI polymorphism
                                                                                                                                                                                                                                                       26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                          ADP08769 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BIOM-)
                                                                                                                                                                                                                                                                                                          ADP08769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lee Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BIOMEDIAB CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GAGTAGCTGGGACTACAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GAGTAGCTGGGATTACAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            be applied as a general screening test
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Baik S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 C; 9 G; 5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kin
Kin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 17.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ŧ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Han
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    S
                                                                                                                               human; platelet PCR; primer; SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Se+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21;
                                                                                                                                                          glycoprotein VI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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WO2004047767-A2 Homo sapiens

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ADO56549
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel method for identifying a subject at risk of breast cancer which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a risk of breast cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Roth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identifying a subject at risk of breast cancer by detecting the presence or absence of one or more nucleotide polymorphic variations, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-NOV-2002; 2002US-0429136P.
24-JUL-2003; 2003US-0490234P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-JUN-2004
                                                                                                                                                                                                                                                                      Human cyclin-dependent kinase 10, CDK10 proximal
                                                                                                                                                                                                                                                                                                 12-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GPIV;GPVI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 3; Page 84; 286pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-NOV-2003; 2003WO-US037966
          WPI; 2004-411721/38
                                                                                  06-NOV-2002; 2002US-0424475P.
23-JUL-2003; 2003US-0489703P.
                                                                                                                                                27-MAY-2004
                                                                                                                                                                        WO2004044164-A2
                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                        single
                                                                                                                                                                                                                                      melanoma associated
                                                                                                                                                                                                                                               gene therapy; human; ss; melanoma;
                                                                                                                                                                                                                                                                                                                                                 ADO56549 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      as well
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-441082/41.
                                                           (SEQU-)
                                                                                                                      06-NOV-2003; 2003WO-US035879
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SEQU-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   well as therapeutic and prophylactic treatments that specifically reget breast cancer, such as gene therapy. The current sequence is that an Extend primer of the invention which was used to genotype single clectide polymorphisms within human glycoprotein VI (platelet) (GP6; IV;GPVI) DNA which is located at chromosomal position 19q13.4.
                                  RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                      869
                                                                                                                                                                                                                                                                                                                                                                                                              19
                                                                                                                                                                                                                       ma associated polymorphic variation; SNP;
nucleotide polymorphism; cyclin-dependent kinase 10; CDK10; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             18;
                                                           SEQUENOM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENOM INC
                                                                                                                                                                                                                                                                                                                                                                                                                            GTTCAAGTTATTCTCCTGC 716
                                                                                                                                                                                                                                                                                                                                                                                                               GTTCAAGTGATTCTCCTGC 1
                                  Nelson MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nelson MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                preventing and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 7 A; 4 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                 (first
                                                           INC
                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.8%;
94.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Braun A,
                                  Braun
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                 Þ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treating breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kammerer SM,
                                  Kammerer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17.4;
No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
.5e+03;
                                    ĸ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Reneland
                                                                                                                                                                                                                                                                          SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                        probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ₽
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                                                                                                                                                                                                                                                                          #74
                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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RESULT 1147
AD056979
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a method of identifying a subject at risk of melanoma comprising detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a subject. Preventing melanoma in a subject comprises detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a subject; and administering a melanoma preventative to a subject in need thereof based upon the presence or absence of the one or more polymorphic variations in the nucleic acid sample. The preventative reduces ultraviolet (UV) light exposure to the subject. The methods, nucleic acids, proteins, and compositions are useful for treating melanoma. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identifying a subject at risk of melanoma, useful for treating melanoma, comprises detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a
                                                                                                                                                                                                                                                                                                                                                                                    melanoma associated polymorphic variation;
single nucleotide polymorphism; CARK; FPGT;
cardiac ankyrin repeat kinase; fucose-1-pho
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 4 A; 2 C; 8
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                                                                              comprises o
                                                                                                                                    WPI;
                                                                                                                                                               Roth
                                                                                                                                                                                                                   06-NOV-2002;
23-JUL-2003;
                                                                                                                                                                                                                                                             06-NOV-2003; 2003WO-US035879
                                                                                                                                                                                                                                                                                                                 WO2004044164-A2
                                                                                                                                                                                                                                                                                                                                                                       probe
                                                                                                                                                                                                                                                                                                                                                                                   cardiac ankyrin
                                                                                                                                                                                                                                                                                                                                                                                                                            gene therapy; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human CARK/FPGT proximal SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AD056979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            represents a human cyclin-dependent kinase 10, CDK10, proximal SNP probe
                                      Example 7; Page 121; 295pp; English
                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                           Identifying a subject at risk of melanoma, useful for treating melanoma, comprises detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a
                                                                                                                                                                                        (SEQU-)
                                                                                                                                                               RB,
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17; Conserv
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                                                                                                                                                                                        SEQUENOM INC
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2003US-0489703P.
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                                                                                                                                                                                                                                                                                                                                                                                                                             ss; melanoma;
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                       probe
                                                                                                                                                                                                                                                                                                                                                                                    fucose-1-phosphate
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The invention relates to a method of identifying a sub melanoma comprising detecting the presence or absence

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RESULT 1148
ADO56537/c
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The invention relates to a method of identifying a subject at risk of melanoma comprising detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a subject preventing melanoma in a subject comprises detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a subject; and administering a melanoma preventative to a subject in need thereof based upon the presence or absence of the one or more polymorphic variations in the nucleic acid sample. The preventative reduces ultraviolet (UV) light exposure to the subject. The methods, nucleic acids, proteins, and compositions are useful for treating melanoma. The present sequence represents a human cyclin-dependent kinase 10, CDK10, proximal SNP probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         melanoma associated polymorphic variation; SNP; single nucleotide polymorphism; cyclin-dependent kinase 10; CDK10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying a subject at risk of melanoma, useful for treating melanoma comprises detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-411721/38.
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23-JUL-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene therapy; human; ss; melanoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human cyclin-dependent kinase 10,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-AUG-2004
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Best Local
                                                                                                 The generic sequence of a primer set designated Alu-2. The primer set was based on bases 240-58 of the 3' end of a 300 bp Alu segment. The primers of the set have an identical sequence to the Alu consensus sequence. Thus priming with the Alu-1 set directs synthesis towards the 3' end (i.e. away from the middle) of the Alu segment. Since the primer set is designed to bind close to the edge of an Alu segment, amplification with these primers will reduce the amount of Alu segment sequence and increase the amount of specific chromosomal DNA present required for probe production. The primer set is useful in the production of chromosomal specific probes e.g for the detection of chromosomal presents such as a probe to detect chronic myelogenous leukemia characterised by the philadelphia chromosome, arising from a reciprocal translocation t(9,22) (q34,q11). (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Primer; PCR; amplification; primer set; probe; Alu sequence; Alu repeat; Alu consensus sequence; chromosome; breakpoint; rearrangement; chronic myelogenous leukemia; Philadelphia chromosome; translocation; ss
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10-AUG-1995
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                                                                                                                                                                                                                                                                                         DNA probe specific for Human chromosome region bcr/abl rearrangement in interphase nuclei.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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                                                                                                                                                                                                                                                                                                                                                    Siciliano MJ,
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RESULT 1150
ABX93649
PXX
                                                   RESULT 1151
ABX95025
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                                                                                                                                                                                                                                                                                                            which the primer binds specifically to the 5'-terminus of an Alu-specific primer, in which the primer binds specifically to the 5'-terminus of an Alu sequence and extends from 3' to 5' of the Alu sequence, or specifically to the 3'-terminus of an Alu sequence and extends from 5' to 3' of the Alu sequence. Also included is a method for preparing genome chips, comprising: (a) obtaining a polynucleotide product by performing the PCR amplification; and (b) spotting the polynucleotide product onto the chip substrate to form the gene chip. The method is used for eliminating a repeat sequence in a genome, which is useful for preparing genome chips from artificial chromosomes for use in diagnosis of genetic diseases, pre-labour diagnosis by screening genetic diseases in pregnant women, tumour typing, diagnosis and prognosis tests, and studying the damaging effects of radioactive rays and other environmental factors on humans. The method allows genome chips to be produced with elimination of Alu repeat sequences and enhanced accuracy by effectively reducing non-specific background signals during hybridisation. The present sequence is an Alu
                                                                                                                                                                                          Matches
                                                                                                                                                                                                         Query Match
Best Local (
ABX95025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; ss; PCR; primer; Alu repeat sequence; artificial chromosome; genome chip; genetic disease; pre-labour diagnosis; tumour typing;
                                  ABX95025 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                              Sequence 20 BP; 5 A; 6 C; 3 G; 3 T; 0 U; 3 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-268207/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-JUL-2001; 2001WO-CN001208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-JUL-2001; 2001WO-CN001208.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 5; Page 8; 18pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            radioactive ray damage; environmental damage.
                                                                                                                                                                                                                                                                                             sequence-specific PCR primer for performing the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to DNA Amplification by polymerase chain reaction (PCR), comprising an artificial chromosome or a large DNA fragment of 50-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eliminating genomic repeat sequences, useful for preparing genome chips from artificial chromosomes for use in diagnosis of e.g. genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYHK-) UNIV HONG KONG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sapiens
                                                                                                                                         871 TTACAGGCGTGAGCCACCAC 890
                                                                                                                                                                                          16;
                                                                                                                      Н
                                                                                                                                                                                                           Similarity
                                                                                                                      TTÁCAGGYRTCAGCCÁCYÁC 20
                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                         1.7%;
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                                                                                                                                                                                                         Score 17.2; DB 1;
Pred. No. 1.5e+03;
                                                                                                                                                                                                                         1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              CC reaction (PCR) is by using an artificial chromosome or a large DNA CC fragment of 50-5000 base pairs in length as template and an Alu-specific CC primer. Also included is a method for preparing a fluorescence-labelling CC probe comprising obtaining a polymucleotide product by performing the PCR caption and fluorescence-labelling the polymucleotide product to performing the PCR complification and fluorescence-labelling the polymucleotide product to CC give the probe. The method is useful for eliminating a repeat sequence in CC a genome, which is applicable in preparing genome chips from artificial CC chromosome for use in diagnosis of genetic diseases, pre-labour diagnosis CC by screening genetic diseases in pregnant women, tumour typing, diagnosis CC and prognosis tests and studying damages of radioactive rays and other CC environmental factors on humans. With this method, RISH (fluorescence incomplete sequence and enhanced accuracy by effectively reducing non-cCC specific background signal during hybridisation. The present sequence CCC represents the human Alu specific PCR primer Alu-N1
                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel method for eliminating repeat sequence in genome, applicable in preparing FISH (fluorescence in-site hybridization) probes from artificial chromosome for use in diagnosis of e.g. genetic diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; ss; PCR; primer; Alu; repeat sequence; fluorescence-labelling; genome chip; pre-labour diagnosis; tumour typing; radioactive ray dam
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003014385-A1
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                 Human; PLAG1; tumourigenesis gene; T-gene; PLAG2; CTNNB1; antibody; benign tumour; malignant tumour; leukaemia; lymphoma; cancer; inhik PCR; amplification; primer; ss.
                                                                                       Nucleotide
                                                                                                                                                                                          AAV29284 standard; cDNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a method of amplification by polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 5; Page 8; 18pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-248303/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-JUL-2001; 2001WO-CN001209.
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                                                                                                                        21-AUG-1998
                                                                                                                                                          AAV29284;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYHK-) UNIV HONG KONG.
                                                                                                                                                                                                                                                                                                      871 TTACAGGCGTGAGCCACCAC 890
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                                                                                      sequence of PCR primer
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                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                 80.0%;
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                                                                                                                                                                                                                                                                                                                                             3;
                                                                                                                                                                                            ВP
                                                                                                                                                                                                                                                                                                                                                                                                                   G; 3 T; 0 U; 3
                                                                                                                                                                                                                                                                                                                                                 Score 17.2; DB 1; Length 20; Pred. No. 1.5e+03; Mismatches 1; Indels
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lymphoma; cancer; inhibition;

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RESULT 1153
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XX AAA2286
XX Integri
XX Integri
XX Human;
KW integri
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This is the nucleotide sequence of the PCR primer Pl used for amplification in the method of the invention, which involves isolation of the tumourigenesis genes (T-gene), in the form of PLAG1, PLAG2, and CTNNB1 genes. Their proteins can be used as a starting point for preparing antibodies for clinically/medically diagnosing cells having a conphysiological proliferative capacity as compared to wild type cells, where the former cells are selected from both benign and malignant tumours, as well as leukaemia and lymphomas. Derivatives of the T-gene care also used in the diagnosis and preparation of therapeutical compositions for the treatment of cancers, such as nucleic acid derivatives, and antibodies. The T-gene may be uses as a starting point for designing suitable expression-modulating compounds or techniques for the treatment of non-physiological proliferation phenomena in humans or animals. Expression inhibitors of the T-gene can be used in the treatment of animals.
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Best Local :
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                                                                   Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Van
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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                                                                                                                                                                                                                                Integrin subunit beta 3 substrate sequence SEQ ID NO:6087.
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17; Conserv
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GOETEBORGS HOLDINGBOLAGET
                                                                                                                                                                                                                                                                  (first entry)
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                                                          syndrome;
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Pred. No.
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                                                                                                                                                                                                                                                                     RESULT 1154
AAA22744
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 13
             Human; aryl hydrocarbon nuclear transport; ARNT; TIB-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiintlammatory; antiarthritic; antipociatic, ARND; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes enzymatic nucleic cleaving activity, which specifically cleave RNA en hydrocarbon nuclear transporter (ARNI) gene, an ir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9950403-A2
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                                                                                                                                                                              19-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                          Integrin subunit beta 3 substrate sequence SEQ ID NO:5970
                                                                                                                                                                                                                  AAA22744;
                                                                                                                                                                                                                                                      AAA22744 standard; RNA; 17
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Pred. No. 1.4e+03;
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psoriasis;

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA23422 represent their corresponding target sequences. The ribozymes the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARMD), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, gooriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, pot-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 3 A; 0 C; 3 G; 0 T; 11 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of an mRNA encoding
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Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes enzymatic nucleic acid molecules with RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pavco PA,
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                                                                                                                                                                                                                                                         AAA22747 standard; RNA; 17
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                                                                                   Integrin subunit beta 3 substrate sequence SEQ ID NO:5973.
                                                                                                                                            19-JUN-2000
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                                                                                                                                            (first entry)
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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RESULT 1156
AAA22759
ID AAA2275
XX
AC AAA2275
XX
AC AAA2275
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DT 19-JUN-

AAA22759 standard; RNA; 17

ВP

19-JUN-2000 AAA22759;

(first entry)

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773

TGTATTTTTAGTAGAGA 789 UGUAUUUUUAGUAGAGA

Matches

9;

Conservative

52.9%;

Score 17; DB 1; Pred. No. 1.4e+03;

Length 17;

Mismatches

0

Gaps

0

Query Match Best Local Similarity

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The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl charges an integrin alpha 6 subunit gene, an integrin subunit beta 3 cg gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, cand AAA17168 to AAA17680 and AAA17623 to AAA17684 represent their corresponding target sequences; AAA17685 to AAA18385 and AAA19087 to AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA19086 cand AAA19155 to AAA19222 represent their corresponding target sequences; AAA19154 represent ribozyme and AAA21595 represent ribozyme corresponding target sequences; CAAA19159 to AAA21681 and AAA21591 to AAA21595 to AAA21500 and CRAA21595 to AAA21687 and AAA21687 to AAA21687 and AAA22475 and AAA22363 to AAA21687 to AAA21687 and AAA22475 and AAA22363 to AAA23342 represent ribozyme sequence for integrin subunit beta 3, and AAA22365 to AAA233343 represent sequence candot integrin subunit beta 3, and AAA22365 to AAA233343 represent sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiotibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; ss. kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss.
                                                                                                      AAA23422 represent their corresponding target sequences. The ribozymes of the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARND), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, psoriasis, vertuca vulgaris, angiofibroma of tuberous sclerosis, pot-wine stains, Sturge Weber angiotome vicinal Transmission and arthribar-pade weber.
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Sequence 17 BP;
                                                                 syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 54; Page 240; 305pp; English.
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                                              subunit alpha-6, or
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5 A; 0 C; 4 G; 0 T; 8 U; 0 Other;
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                                              subunit beta-3
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RESULT 1157
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Integrin subunit beta 3 substrate sequence SEQ ID NO:5985

Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiintlammatory; antiarthritic; antipsoriatic; ARND; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss

Homo sapiens

07-OCT-1999.

24-MAR-1999; 99WO-US006507

27-MAR-1998; 98US-0079678P

(RIBO-) RIBOZYME PHARM INC

Roberts E, Jarvis 'n Coeshott C, Mcswiggen

WPI; 1999-591315/50

ribozymes for modulating the synthesis,  $\ensuremath{\mathsf{mRNA}}$  encoding an angiogenic factors. expression and/or stability

Claim 54; Page 240; 305pp; English

The present invention describes enzymatic nucleic acid molecules with RNA Cc cleaving activity, which specifically cleave RNA encoded by an aryl Cc hydrocarbon nuclear transporter (ARNI) gene, an integrin subunit beta 3 CC gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to CC AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, CC and AAA17168 to AAA17563 and AAA17625 to AAA17684 represent their Cc corresponding target sequences; AAA17685 to AAA18385 and AAA19087 to CC AAA19157 to AAA19158 co AAA19158 to AAA19086 cc and AAA19155 to AAA12037 to AAA17685 to AAA19223 to AAA21681 and AAA21501 to AAA21595 represent traces; CC AAA21922 to AAA21688 represent their corresponding target sequences; CC AAA21896 to AAA21688 represent their corresponding target sequences; CC AAA21896 to AAA21688 represent their corresponding target sequences; CC AAA21892 to AAA22475 and AAA22361 to AAA2342 represent ribozyme sequence for integrin subunit beta 3, and AAA22476 to AAA23362, AAA23343 to CC aAA23422 represent their corresponding target sequences; CC the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding anglogenic factor, especially ARNT, CC integrin subunit beta 3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related CC macular degeneration (ARMD), inflammation, and arthritis, as well as especially come, kippel-Trenaunay-Weber syndrome, stains, Sturge Weber come, subcome, sub

Sequence 17 BP; 4 A; 7 C; 3 G; 0 T; 3 U; 0 Other;

Query Match Best Local Matches Similarity 82.4%; Pred. Score 17; Mismatches No. 1.4e+03; DB 1; 0 Length 17; Indels ٥, Gaps

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Ś 378 CTCAGCCTCCCAAAGTG 394

> AAA22860 standard; RNA; 17 19-JUN-2000 (first entry) BP.

Integrin subunit beta 3 substrate sequence SEQ ID NO:6086

Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiintlammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss

Homo sapiens

WO9950403-A2

07-OCT-1999.

24-MAR-1999; 99WO-US006507

27-MAR-1998; 98US-0079678P

(RIBO-) RIBOZYME PHARM INC

Pavco PA, Roberts E, Jarvis 'n Coeshott C, Mcswiggen JA;

WPI; 1999-591315/50

Novel ribozymes for modulating the synthesis, of an mRNA encoding an angiogenic factors. expression and/or stability

Claim 54; Page 247; 305pp; English

CC cleaving activity, which specifically cleave RNA encoded by an aryl activity and integrin alpha 6 subunit gene, an integrin subunit beta 3 cc gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to CC AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, CC corresponding target sequences; AAA1685 to AAA18365 and AAA19087 to CC corresponding target sequences; AAA1685 to AAA18365 and AAA19087 to CC carresponding target sequences for Tie-2, and AAA19186 to AAA19086 cc AAA19154 represent ribozyme sequences for Tie-2, and AAA19186 to AAA19086 cc AAA19154 represent their corresponding target sequences; CC AAA19123 to AAA20361 and AAA218501 to AAA21855 represent ribozyme cc sequences for integrin alpha 6 subunit, and AAA20362 to AAA21500 and CC AAA21596 to AAA21688 represent their corresponding target sequences; CC AAA21689 to AAA22475 and AAA23273 to AAA2342 represent ribozyme sequence for integrin subunit beta 3, and AAA22476 to AAA23232, AAA233343 to CC AAA23422 represent their corresponding target sequences of CC integrin subunit beta 3, and AAA22476 to AAA23262, AAA33343 to CC attegrin subunit beta 3, and AAA22476 to AAA23262, AAA33343 to CC attegrin subunit beta 3, integrin subunit alpha-6, or Tie-2. They are CC especially used to treat cancer, diabetic retinopathy, age related CC macular degeneration (ARMD), inflammation, and arthritis, as well as covascular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, cC angiofibroma of tuberous sclerosis, pot-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu and other syndromes and diseases related to the levels of ARN integrin subunit alpha-6, or integrin subunit beta-3 The present invention describes enzymatic nucleic acid molecules with RNA ndu syndrome, ARNT, Tie-2,

Sequence 17 BP; 3 A; 3 C; 7 <u>ဂ</u> 0 T; 4 U; 0 Other;

Matches Query Match Local Similarity 1.7%; Score 17; DB Pred. No. 1.46 4; Mismatches 1.4e+03; DB 1; Length 17; Indels 0 Gaps

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394 GCTGGGATTACAGGCGT

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GCUGGGAUUACAGGCGU 17

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RESULT 1158
                                                                                                        CC cleaving activity, which specifically cleave RNA encoded by an aryl C hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 CC gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to CC AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, CC and AAA17168 to AAA17663 and AAA17684 represent their CC corresponding target sequences; AAA17685 to AAA18385 and AAA19087 to CC AAA19154 represent ribozyme sequences for Tie-2, and AAA18986 to AAA19086 CC and AAA19155 to AAA12222 represent their corresponding target sequences; CC AAA19154 represent ribozyme sequences; CC AAA19154 represent ribozyme sequences; CC AAA19223 to AAA21861 and AAA21501 to AAA21595 represent ribozyme CC sequences for integrin alpha 6 subunit, and AAA22325 to AAA21500 and AAA19223 to AAA21688 represent their corresponding target sequences; CC AAA21596 to AAA21688 represent their corresponding target sequences; CC AAA21596 to AAA22475 and AAA23263 to AAA2342 represent ribozyme sequence for integrin subunit beta 3, and AAA22476 to AAA23343 to CC AAA21689 to AAA22475 and AAA22476 to AAA22362, AAA23343 to CC AAA23422 represent their corresponding target sequences. The ribozymes of CC AAA23422 represent their corresponding target sequences of CC CAA23422 represent their corresponding target sequences. The ribozymes of CC atability of an mRNA encoding angiogenic factor, especially ARNT, CC integrin subunit beta 3, integrin subunit alpha-6, or Tie-2. They are CC especially used to treat cancer, diabetic retinopathy, age related CC macular degeneration (ARMD), inflammation, and arthritis, as well as covascular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, CC avadical subunit subarous sclerosis, pot-wine stains, Sturge Weber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 54; Page 239; 305pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pavco PA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes enzymatic nucleic acid molecules with RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC.
                                                                    drome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, other syndromes and diseases related to the levels of ARNT, Tie-2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mRNA encoding an angiogenic factors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ribozymes for modulating the synthesis, expression and/or stability
                                            subunit alpha-6, or integrin subunit beta-3
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Query Match Sequence

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                   The present invention describes enzymatic nucleic acid molecules with RNA CC cleaving activity, which specifically cleave RNA encoded by an aryl CC hydrocarbon nuclear transporter (ARNT) gene, an integrin alpha 6 subunit gene, or a file-2 gene. AAA16775 to CC AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, CC and AAA17168 to AAA17560 and AAA17685 to AAA16895 and AAA19087 to CC CAA19154 represent ribozyme sequences; AAA17685 to AAA18385 and AAA19087 to CC AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA19086 CC AAA19154 represent ribozyme sequences; CC AAA19155 to AAA19222 represent their corresponding target sequences; CC AAA19155 to AAA20361 and AAA21551 to AAA21595 represent ribozyme sequences; CC AAA19153 to AAA20361 and AAA21551 to AAA21595 represent ribozyme (CC AAA21596 to AAA21688 represent their corresponding target sequences; CC AAA21596 to AAA22475 and AAA23243 to AAA21500 and CC AAA21596 to AAA22475 and AAA23253 to AAA2342 represent ribozyme sequence (CC for integrin subunit beta 3, and AAA22476 to AAA23343 to CC CAA23422 represent their corresponding target sequences. The ribozymes of the invention are used for modulating the synthesis, expression and/or (CC stability of an mRNA encoding angiogenic factor, especially ARNT; CC integrin subunit beta 3, integrin subunit alpha-6, or Tie-2. They are (c) especially used to treat cancer, diabetic retinopathy, age related (CC macular degeneration (ARMD), inflammation, and arthritis, as well as conditors englancema, myopic degeneration, psoriasis, verruca vulgaris, canditors, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu Syndrome, and diseases related to the levels of DNT Tie-2.
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Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis, age related macular degeneration; inflammation; neovascular glaucoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 54; Page 238; 305pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-591315/50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Integrin subunit beta 3 substrate sequence SEQ ID NO:5948.
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   other syndromes
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       and diseases
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   related to the
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Matches 13
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                                                                                      The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNI) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA1675 to AAA17671 and AAA17561 to AAA17622 represent ribozyme sequences for ARNIT, and AAA17168 to AAA17620 and AAA17623 to AAA17644 represent their corresponding target sequences; AAA17685 to AAA18385 and AAA19087 to AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA19086 and AAA19155 to AAA19222 represent their corresponding target sequences; AAA19223 to AAA19223 represent their corresponding target sequences; AAA19223 to AAA20361 and AAA21501 to AAA21595 represent ribozyme sequences for integrin alpha 6 subunit, and AAA20362 to AAA21500 and AAA21596 to AAA21688 represent their corresponding target sequences for integrin subunit beta 3, and AAA23470 represent ribozyme sequence for integrin subunit beta 3, and AAA22476 to AAA23343 to AAA23343 represent ribozyme sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human; aryl hydrocarbon nuclear transport; ARNY; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD;
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AAA23422 represent their corresponding target sequences. The ribozymes the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT, integrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                              mRNA encoding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      neovascular glaucoma, myopic degeneration, psoriasis, verruca vulangiofibroma of tuberous sclerosis, pot-wine stains, Sturge Weber Syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndromes and diseases related to the levels of ARNT, Tintegrin subunit alpha-6, or integrin subunit beta-3
The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, and AAA17168 to AAA17623 to AAA17684 represent their corresponding target sequences; AAA17623 to AAA18385 and AAA19087 to AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA19086 and AAA19122 represent their corresponding target sequences; AAA19223 to AAA20361 and AAA21501 to AAA21595 represent ribozyme sequences for Tie AAA21596 to AAA21598 represent their corresponding target sequences; AAA19223 to AAA20361 and AAA21503 to AAA20362 to AAA21500 and AAA21596 to AAA21488 represent their corresponding target sequences; AAA21596 to AAA221488 represent their corresponding target sequences; AAA21596 to AAA22475 and AAA233343 to AAA23342 represent tribozyme sequences; AAA221689 to AAA22475 and AAA233343 represent tribozyme sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozy hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antiporiatic; dermatological; RNA cleavage; cancer; dlabetic retinopathy; arthrage related macular degeneration; inflammation; neovascular glauc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 4 A; 7 C; 2 G; 0 T; 4 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-OCT-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; aryl hydrocarbon nuclear transport; ARNT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-JUN-2000
                                                                                                                                                                                                                                                                     Claim 54;
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                                                                                                                                                                                                                                                                                                      ribozymes for modulating the synthesis, mRNA encoding an angiogenic factors.
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                     The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA171367 and AAA17631 to AAA17622 represent ribozyme sequences for ARNT, and AAA17168 to AAA17660 and AAA17623 to AAA17644 represent their corresponding target sequences; AAA17685 to AAA18385 and AAA19087 to AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA19086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARND; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; age related macular degeneration; inflammation; neovascular glaucoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Roberts E,
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AAA23422 represent their corresponding target sequences. The ribozymes the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT, integrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARMD), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, pot-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA19223 to AAA20361 and AAA21501 to AAA21595 represent ribozyme sequences for integrin alpha 6 subunit, and AAA20362 to AAA21500 and AAA21596 to AAA21688 represent their corresponding target sequences; AAA21596 to AAA22475 and AAA23263 to AAA23342 represent ribozyme sequence for integrin subunit beta 3, and AAA22476 to AAA23262, AAA23343 to
                                                                                                                                                           integrin subunit alpha-6, or integrin subunit beta-3
17 BP; 4 A; 0 C; 4 G;
0
T; 9 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              O<sub>Ff</sub>
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Matches
                                      Query Match
Best Local
                772
1 UUGUAUUUUUAGUAGAG
                               Similarity
8; Conserv
                TTGTATTTTTAGTAGAG
                                Conservative
                                      1.78;
                788
17
                                9;
                                       Score 17; I
Pred. No. 1.
                                 Mismatches
                            DB 1;
1.4e+03;
0;
                                               Length
                                0
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0

밁 S

Integrin subunit beta 3 substrate sequence SEQ ID NO:5971 19-JUN-2000 AAA22745; AAA22745 standard; (first entry) RNA; 17 ВÞ

ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; arthritis; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; age related macular degeneration; inflammation; andiofibroma; myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; Human; aryl hydrocarbon nuclear transport; Homo sapiens hammerhead Kippel-Trenaunay-Weber integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozy
nammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic,
phthalmologic; antiinflammatory; antiarthritic; antipsoriatic; // syndrome; Osler-Weber-Rendu syndrome; ss ARNT; TIE-2; angiogenesis; ribozyme;

WO9950403-A2

24-MAR-1999; 99WO-US006507

27-MAR-1998; 98US-0079678P

RIBOZYME PHARM

PA, Roberts E, Jarvis H Coeshott Ç Mcswiggen

WPI; 1999-591315/50

of an ribozymes for modulating mRNA encoding an angioger encoding angiogenic the synthesis, expression and/or stability

54; Page 239; 305pp; English.

The present invention describes enzymatic nucleic acid molecules with cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta gene, subunit gene, S S an integrin Tie-2 gene.

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RESULT 1164
AAA22831
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PRINCE PR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IJ
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Best Local S
Matches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiintlammatory; antiarthritic; antipsoriatic; ARND; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; age related macular degeneration; inflammation; neovascular glaucoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        myopic degeneration; psoriasis; verruca vulgaris; angiofibroma;
tuberous sclerosis; pot-wine stain; Sturge Weber syndrome;
Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-JUN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA22831 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9950403-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Integrin subunit beta 3 substrate sequence SEQ ID NO:6057.
                                                                                                                                                                      WPI; 1999-591315/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                      27-MAR-1998;
                                                                                                                                                                                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 41.2
les 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            771 TTTGTATTTTTAGTAGA 787
                                                                                                                                                                                                                                                 PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 UUUGUAUUUUUAGUAGA 17
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                                                                                                                                                                                                                                                 Roberts E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP;
                                                                                                                                                                                                                                                                                                                                                                                      98US-0079678P
                                                                                                                                                                                                                                                                                                                                                                                                                                                             99WO-US006507
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.7%;
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                                                                                                                                                                                                                                                 Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 17; DB 1;
Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                    Coeshott C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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                                                                                                                                                                                                                                                 Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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The present invention describes enzymatic nucleic acid molecules with RNA CC cleaving activity, which specifically cleave RNA encoded by an aryl CC hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 CC gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA17616 to AAA17622 represent ribozyme sequences for ARNT, CC corresponding target sequences; AAA17685 to AAA18385 and AAA19087 to CC AAA19154 represent ribozyme sequences for Tie-2, and AAA19087 to CC AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA19086 cand AAA19155 to AAA12322 represent their corresponding target sequences; AAA17685 to AAA18385 and AAA19086 cand AAA19155 to AAA12321 to AAA20361 and AAA21501 to AAA21355 represent ribozyme sequences for Tie-2, and AAA19086 cand AAA21323 to AAA21588 represent their corresponding target sequences; AAA19086 cand AAA19232 represent their corresponding target sequences; AAA21595 to AAA21595 to AAA22342 represent their corresponding target sequences for integrin subunit beta 3, and AAA23476 to AAA23342 represent ribozyme sequence for integrin subunit beta 3, and AAA22476 to AAA23342 represent their corresponding target sequences. The ribozymes of the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT, CC integrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related corresponding target sequences of tuberous sclerosis, pot-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT.
   Sequence 17
                                                                        integrin subunit alpha-6, or integrin subunit beta-3
BP;
       _
       Þ
       10 C; 2 G; 0 T; 4 U; 0 Other;
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RESULT 1165
AAC87597/c
                                                                                                                                             Matches
                                                                                                                                                        Query Match
Best Local :
16-MAR-2001
                      AAC87597;
                                            AAC87597 standard; DNA; 17
                                                                                                                       536 TCCTGCCTCAGCCTCCC 552
                                                                                                 1 UCCUGCCUCAGCCUCCC 17
                                                                                                                                                        Similarity
                                                                                                                                              Conservative
(first entry)
                                                                                                                                                         76.5%;
                                                                                                                                             4.
                                                                                                                                                         Score 17;
Pred. No.
                                                                                                                                                Mismatches
                                                                                                                                                         1.4e+03;
                                                                                                                                                                    DB 1;
                                                                                                                                                                    Length 17
                                                                                                                                                Indels
                                                                                                                                                <u>,,</u>
                                                                                                                                                 Gaps
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Human; keratinocyte growth factor; KGF; chromosome 9p11; abnormality; cancer; miscarriage; spontaneous abortion; genetic susceptibility; diagnosis; Alu sequence; PCR primer; ss.

Human Alu sequence PCR primer,

CL2.

Homo sapiens

JP2000287684-A

17-OCT-2000.

31-JAN-2000; 2000JP-00022688

05-FEB-1999; 99JP-00028705

(ASAK ) ASAHI BREWERIES TTD

WPI; 2001-065570/08.

The base sequence of 9p11 chromosomal region participating 6 cancer and

3; Page 5; 88pp; Japanese

ribozymes for modulating the synthesis, expression and/or stability mRNA encoding an angiogenic factors.

54; Page 245; 305pp; English

encoding an angiogenic

The invention relates to human chromosomal region 9q11 (AAC87588). Abnormalities in this region of the short arm of chromosome 9 is thought

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ADB04439
ID ADB0443
XX Cytosta
XW Cytosta
XW Zinc fi
XW Chromos
XW Zinc fi
XW Chromos
XW Zinc fi
XW Chromos
XW Advelop
XX Homo sa
XX EP12817
XX O5-FEB-
XX O2-AUG-
PF 30-JUL-
XX NEW Zin
PF MADZ4, M
XX The pre
CC Protein
CC Encoded
CC MDZ7 in m
CC associa
CC MDZ7. M
CC acids a
CC alterat
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               to be associated with miscarriage and cancer, as an ovarian cancer patient with a history of miscarriage was found to have a chromosomal inversion inv(9) (pl1;q13). The 9p11 region contains the gene encoding keratinocyte growth factor (KGF), and the invention also specifically claims the KGF PCR primers AAC87589 and AAC87590 for use in detecting all or part of the KGF gene. The nucleic acid sequences can be used to detect abnormalities in chromosomal region 9p11 and thus give an indication of an individual's risk of developing a 9p11-associated condition. Sequences AAC87596-C87597 represent human Alu sequence PCR primers used in an exemplification fo the invention
                          proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome fsp1.3-22.2, MDZ7 is encoded at chromosome fsp1.2 and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADB04439 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; Chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB04439;
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                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to novel human zinc finger-containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              associated MDZ4, MDZ7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-JUL-2002; 2002EP-00016874.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human MDZ7 scanning oligonucleotide SEQ ID 5425.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2003-423107/40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   with decreased or incr
or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ည
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 2 A; 7 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ID NO 5425; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Y, Nguyen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>.</u>
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constructing microarrays for measuring

gene expression.

Query Match
Best Local Similarity
Matches 17; Conserv

1.7%; ilarity 100.0%; Conservative

0

Score 17; DB; Pred. No. 1.4

DB 1; . 1.4e+03

Length 17

Indels

0

Gaps

0

Sequence 17 BP;

5 A; 0 C; 4 G; 8 T; 0 U; 0 Other;

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RESULT 1167
ADB04442
ID ADB0444
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                                       The present invention relates to novel human zinc finger-containing CC proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 5p21.3-22.2, CC MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome CC or in manufacturing a medicament for treating or preventing a disorder CC associated with decreased or increased expression or activity of MDZ3, CC MDZ4, MDZ7, c.e.g. cancer or developmental disorders. The nucleic CC acids and proteins are also useful for diagnosing or monitoring a disease CC caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic CC acids can also be used as probes to detect and characterize gross CC alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                  Example 8; SEQ ID NO 5428; 103pp; English.
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                               present sequence
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RESULT 1168
ADB04282
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XX Cytosts
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                                                                                                                                                         Query Match
Best Local
                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                                                         Sequence 17 BP; 4 A; 2 C; 4 G; 7 T; 0 U; 0 Other;
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RESULT 1169 ADB04440

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               Human MDZ7 scanning oligonucleotide SEQ
                                               20-NOV-2003
                                                                                                                                                                                                                                                                                                                      Sequence 17
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                                                                               ADB04314;
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                                                                                                                                                                                                                                                                                                                          BP; 4
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                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                          3 G; 10 T; 0
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                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7922.1, MDZ41 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 16p26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 is equences are useful in therapy. Or in manufacturing a medicament for treating or preventing a disorder 2 associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease in caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
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Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-JUL-2002; 2002EP-00016874
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                                                                                                                                   Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD23; MD24; MD27; MD212; chromosome 7 chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 2 A; 3 C; 9 G; 3 T; 0 U; 0 Other;
       EP1281758-A2
                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                              Human MDZ7
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                                                                                                                                                                                                                                                                                                       scanning oligonucleotide SEQ ID 5269.
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Pred. No.
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                                                                                                                                                                       15q26.1;
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RESULT 1172
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PA PR XXX PX XXX XXX PR XXX PR
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Matches 17; Conservative (
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Best Local :
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                                                         02-AUG-2001; 2001US-00922181
                                                                                                                          30-JUL-2002; 2002EP-00016874.
                                                                                                                                                                                                                                                            EP1281758-A2
                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human MDZ7 scanning oligonucleotide SEQ ID
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(AEOM-) AEOMICA INC

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RESULT 1173
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06-JUN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-rheumatic; cancer; AIDS; 88.
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   Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding
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                                                                                                                        WPI; 2003-140484/13
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c; anti-HIV;
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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ra human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are

N-Ras,

treating cand HER2, K-Ras,

Claim

58; Page 98; 185pp; English.

WPI; 2003-140484/13.

(RIBO-) RIBOZYME

PHARM INC

Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encod HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.

encoding

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim
                                                                                                                                                                                                                                                                                                                                                                                                                           Human, ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17
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06-JUN-2001; 2001US-0296249P
10-SEP-2001; 2001US-0318471P
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76.5%;
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         sequence having at least 80% identity, after optimal alignment, with the cuncleotides, a sequence that hybridizes under stringent conditions with the nucleotides, or the complement, or corresponding RNA, of the nucleotides. The nucleotides are used as probes or primers for detecting, identifying, quantifying and/or amplifying nucleic acids, as in vitro sense and antisense sequences, of nucleotides involved in tumour suppression apoptosis and or viral resistance, to produce recombinant polypeptides, and to prepare transgenic animals, as experimental models. The nucleotides (also vectors containing the wectors), the encoded polypeptides and antibodies (Ab) against the polypeptide are useful for prevention and/or treatment of viral infections or diseases characterized by development of tumours or calls degeneration (e.g. Alzheimer's disease or schizophrenia).

Analysis of the expression of the nucleotides can be used for diagnosis and/or prognosis of these diseases. The nucleotides and polypeptides can
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                                                                                                                                                                                                                                                                                    The invention relates to the isolation of 6327 nucleotide sequences, fragments of at least 15 consecutive nucleotides of these nucleotides,
                                                                                                                                                                                                                                                                                                                                                                            useful e.g.
polypeptide
                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related
                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-441574/41.
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virus resistance; transgenic animals; Alzheimer's disease; schizophrenia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic; antiviral; neuroprotective; nootropic; neuroleptic;
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                     least 20 but not more than 1500 consecutive nucleotides of the optineurin correctory promoter appearing as ADE(13890. Also included are the optineurin promoter operably linked to a heterologous nucleic acid, a nucleic acid capable of detecting a single nucleotide polymorphism (SNP) in the optineurin promoter promoter, a host cell comprising the promoter operably linked to a heterologous sequence, diagnosing or prognosing glaucoma in a sample of heterologous sequence, diagnosing of the promoter plantom a cell or bodily fluid (comprising detecting a polymorphism in a promoter region of the optineurin gene, associated with a glaucoma complementype), detecting a SNP sequence variation in a sample containing DNA, detecting a sequence variation in a sample containing DNA, detecting the presence of an optineurin promoter sequence variation in a sample containing DNA, determining the presence or increased susceptibility to glaucoma or to a progressive ocular hypertensive consceptibility to glaucoma or to a progressive ocular hypertensive conspicuation reaction primers that direct amplification of a selected amplification reaction primers that direct amplification of a selected concleic acid region containing the variation within the optineurin comprising obtaining a sample containing the DNA) and detecting a polymorphism (comprising capable of detecting a SNA) and detecting a polymorphism comprise and cide canable of detecting a SNA placeted within an optineurin promoter and amid in the soul of the propersion of the service of the propersion 
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glaucoma related disorder; motif; repeat element; regulatory region.
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   obtaining a sample co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 11; SEQ ID NO 354; 159pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Raymond V,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       optineurin promoter to
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RAYMOND V.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          glaucoma and related
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located within an optineurin

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RESULT 1177
ADH59606/c
 Matches 17;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              detecting the polymorphism). The inv
prognose glaucoma and also to treat
present sequence is an optineurin pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 11 A; 3 C; 0 G; 3 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             putative regulatory region.
                                                comprising 5-50 probes), contacting the sample with the one or more detectable nucleic acid probes, and determining the target genomic nucleic acid of the sample by determining the hydridization of the one or more detectable nucleic acid probes to the target genomic nucleic acid of the sample. The genomic nucleic acid is contained in a fixed tissue or a cell, and the sample is metaphases spreads, interphase nucleic or nucleic found in paraffin embedded tissue material or frozen tissue sections. The probe is also useful in comparing a sample of genomic nucleic acid with that of a control sample using a genomic nucleic acid array. The method comprises treating a sample of genomic nucleic acid and control genomic nucleic acid, which are differentially labelled, the
                                                                                                                                                                                                                                                                                                             The present sequence represents a non-nucleotide probe. The probe is useful for suppressing the binding of one or more detectable nucleic probes, that are greater than 100 base pairs and that have been deriven.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Non-nucleotide probe for suppressing binding of detectable nucleic acid probes to undesired sequences, has aggregate nucleobase sequence homologous to randomly distributed repeat sequence of genomic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-SEP-2001; 2001US-0324499P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              non-nucleotide probe; Bacterial Artificial Chromosome clone; BAC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Non-nucleotide probe of the invention #10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADH59606 standard; DNA; 17
array or both the sample and control genomic nucleic acid and the arrawith the mixture of the probe under suitable hybridization conditions contacting the array with treated mixture of sample and control genome
                                                                                                                                                                                                                                                         probes, that are greater than 100 base pairs and that have been derived from genomic nucleic acid, to one or more undestred sequences in an assifor determining trayet genomic nucleic acid of a sample. The method comprises contacting the sample with the mixture of probes (preferably
                                                                                                                                                                                                                                                                                                                                                                                          Claim 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kirtsen NV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BOST-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            770
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BOSTON PROBES INC
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                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 12; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hyldig-Nielsen JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention is used to diagnose and o treat glaucoma related disorders. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Williams BF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  promoter motif,
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     control genomic
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                                                                                                               Best
                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                             nucleic acid under suitable hybridization conditions, and comparing the intensities of the signals from the differential labels of the array to that caused by hybridization of the probes to genomic nucleic acid, thus determining one or more variations in copy numbers of sequences in the sample as compared with the relative copy numbers of substantially identical sequences in the control. The hybridization of the genomic array is determined using an intercalating dye or a detectable antibody, or its fragment, that is specific for a nucleic acid/nucleic acid hybrid. The sample of genomic nucleic acid to be tested and the reference of nucleic acid are labelled with detectable moiety such that hybridization for the control of the sample of genomic nucleic acid to be tested and the reference of nucleic acid are labelled with detectable moiety such that hybridization
                                                                                                                                                                                                                                                       of the genomic array is determined by determining the presence, absence, amount or location of the detectable label on the one or more genomic arrays. The genomic array comprises nucleic acid that is prepared from Bacterial Artificial Chromosome (BAC) clones. The present sequence
                                                                                                                                                                                   Sequence 17
                                                                                                                                                                                                                                  represents a non-nucleotide probe of the invention.
                                                                                                               Local
                                              967
17
                                                                                            l Similarity
17; Conserv
                                              ATCTCGGCTCACTGCAA 983
  ATCTCGGCTCACTGCAA
                                                                                                                                                                                        BP; 4
                                                                                            Conservative
                                                                                                                                                                                      A; 3 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                     100.0%; -
    <u>.</u>
                                                                                                                  Score 17;
Pred. No.
                                                                                                 Mismatches
                                                                                                                       No.
                                                                                                                                               멂
                                                                                                                    .4e+03;
                                                                                                                                          Length 17;
                                                                                                 Indels
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RESULT 1178
ADH59604/c
non-nucleotide probe; Bacterial Artificial Chromosome clone; BAC;
                                                                                                                   Synthetic
                                                                                                                                      Non-nucleotide
                                                                                                                                             25-MAR-2004
                                                                                                                                                    ADH59604;
                                                                                                                                                           ADH59604 standard; DNA; 17
                                                                                                                                             (first entry)
                                                                                                                                      probe of the invention
                                                                                                                                      #8.
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88;

WO2003027328-A2

03-APR-2003.

24-SEP-2002; 2002WO-US030573.

24-SEP-2001; 2001US-0324499P

(DAKO-) DAKOCYTOMATION DENMARK BOSTON PROBES INC

AS

Kirtsen Š Hyldig-Nielsen JJ, Williams BF

WPI;

Non-nucleotide probe for suppressing binding of detectable nucleic acid probes to undesired sequences, has aggregate nucleobase sequence homologous to randomly distributed repeat sequence of genomic nucleic homologous esired sequences, has randomly distributed

Claim 10; SEQ ID NO 10; 103pp; English

The present sequence represents a non-nucleotide probe. The probe is useful for suppressing the binding of one or more detectable nucleic probes, that are greater than 100 base pairs and that have been derive from genomic nucleic acid, to one or more undesired sequences in an a for determining target genomic nucleic acid of a sample. The method comprises contacting the sample with the mixture of probes (preferable) comprising 5-50 probes), contacting the sample with the one detectable nucleic acid probes, and determining the target of nucleic acid of the sample by determining the hybridization the one or more genom (preferably derived one аввау acid or P

Claim 10; SEQ ID NO 22; 103pp; English

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RESULT 1179
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 17
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Non-nucleotide probe for suppressing binding of detectable nucleic acid probes to undesired sequences, has aggregate nucleobase sequence homologous to randomly distributed repeat sequence of genomic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Non-nucleotide probe of the invention #20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-SEP-2002; 2002WO-US030573
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADH59616 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      arrays. The genomic array comprises nucleic acid that is prepared from Bacterial Artificial Chromosome (BAC) clones. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The method comprises treating a sample of genomic nucleic acid a control genomic nucleic acid, which are differentially labelled,
                                                                                                                                                                                                                WPI; 2003-421160/39
                                                                                                                                                                                                                                                                                                       Kirtsen NV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-SEP-2001; 2001US-0324499P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  non-nucleotide probe; Bacterial Artificial Chromosome clone; BAC; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         represents a non-nucleotide probe of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                       (BOST-) BOSTON PROBES INC. (DAKO-) DAKOCYTOMATION DENMARK AS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                536 TCCTGCCTCAGCCTCCC 552
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 4 A; 2 C; 10 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                       Hyldig-Nielsen JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.7%;
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. 1.,
                                                                                                                                                                                                                                                                                                            Williams BF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.4e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        acid and
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CC comprises contracting the sample with the mixture of probes (preferably comprises contracting the sample with the mixture of probes) preferably comprising 5-50 probes), contacting the sample with the one or more detectable mucleic acid probes, and determining the target genomic cucleic acid of the sample by determining the hybridization of the one or comore detectable mucleic acid probes to the target genomic nucleic acid of the sample. The genomic nucleic acid is contained in a fixed tissue or a color of the one or color of the sample. The genomic nucleic acid is contained in a fixed tissue or a color of the sample is metaphase spreads, interphase nucleic or nucleic color of a control sample using a genomic nucleic acid with comprises treating a sample of genomic nucleic acid and control genomic nucleic acid and the array or both the sample and control genomic nucleic acid and the array or both the mixture of the probe under suitable hybridization conditions, contacting the array with treated mixture of sample and control genomic nucleic acid under suitable hybridization conditions, and comparing the contacting one or more variations in copy numbers of sequences in the carray is determined using an intercalating dye or a detectable antibody, correspondic array is determined using an intercalating dye or a detectable antibody, correspondic array is determined using an intercalating dye or a detectable antibody, or its fragment, that is specific for a nucleic acid/nucleic acid hybrid. The sample of genomic array is determined by determining the presence, absence, amount or location of the detectable moiety such that hybridization continued in the order of the detectable and the order genomic anon-nucleotide with detectable and the reference of anon-nucleotide probe of the hybridization or or more genomic control array is determined by determining 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  useful for suppressing the binding of one or more detectable nucleic probes, that are greater than 100 base pairs and that have been der from genomic nucleic acid, to one or more undesired sequences in an fordetermining target genomic nucleic acid of a sample. The method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents a non-nucleotide probe. The probe is useful for suppressing the binding of one or more detectable nucleic
represents a non-nucleotide probe of the invention.
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밁
                                 Query Match
Best Local Similarity
Matches 17; Conserv
                                 Sequence 17 BP; 1 A; 10 C; 2 G; 4 T; 0 U; 0 Other;
     536 TCCTGCCTCAGCCTCCC 552
ш
TCCTGCCTCAGCCTCCC 17
                Conservative
                    100.0%;
                ,
,
                    Score 17;
Pred. No.
                 Mismatches
                        DB 1;
                    1.4e+03;
                        Length 17;
                Indels
                <u>.</u>
                Gaps
                0
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RESULT 1180
ADH59618
ID ADH5961
Non-nucleotide probe of the invention #22.
                                                                                                                                                                   ADH59618;
               24-SEP-2001; 2001US-0324499P
                                24-SEP-2002; 2002WO-US030573.
                                                                    WO2003027328-A2
                                                                                     Synthetic
                                                                                                                                                  25-MAR-2004 (first entry)
                                                                                                                                                                                    ADH59618 standard; DNA; 17
                                                                                                        probe.
                                                                                                               non-nucleotide probe; Bacterial Artificial Chromosome clone; BAC;
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(BOST-) BOSTON PROBES INC

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ACC51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    countrol genomic nucleic acid, which are differentially resolved, or both the sample and control genomic nucleic acid and the array contracting the array with treated mixture of sample and control genomic nucleic acid under suitable hybridization conditions, contacting the array with treated mixture of sample and control genomic nucleic acid under suitable hybridization conditions, and comparing the contensities of the signals from the differential labels of the array to that caused by hybridization of the probes to genomic nucleic acid, thus contensities of the array to contensity one or more variations in copy numbers of sequences in the contensity of substantially contensity of array is determined using an intercalating dye or a detectable antibody, cor its fragment, that is specific for a nucleic acid/nucleic acid hybridical corresponding array is determined by determining the presence, absence, comunit or location of the detectable moisty such that hybridization of the genomic array is determined by determining the presence, absence, comunit or location of the detectable label on the one or more genomic carrays. The genomic array comprises nucleic acid that is prepared from contensing an artificial Chromosome (BAC) clones. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        more detectable nucleic acid probes to the target genomic nucleic acid of the sample. The genomic nucleic acid is contained in a fixed tissue or a cell, and the sample is metaphase spreads, interphase nucleic or nucleic found in paraffin embedded tissue material or frozen tissue sections. The probe is also useful in comparing a sample of genomic nucleic acid with that of a control sample using a genomic nucleic acid reference array. The method comprises treating a sample of genomic nucleic acid and control genomic nucleic acid, which are differentially labelled, the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          from genomic nucleic acid, to one or more undesired sequences in an assay for determining target genomic nucleic acid of a sample. The method comprises contacting the sample with the mixture of probes (preferably comprising 5-50 probes), contacting the sample with the one or more detectable nucleic acid probes, and determining the target genomic nucleic acid of the sample by determining the hybridization of the one or nucleic acid of the sample by determining the hybridization of the one or nucleic acid of the sample by determining the hybridization of the one or nucleic acid of the sample by determining the hybridization of the one or nucleic acid of the sample by determining the hybridization of the one or nucleic acid of the sample by determining the hybridization of the one or nucleic acid of the sample by determining the hybridization of the one or nucleic acid of the sample by determining the hybridization of the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid probes acid acid probes with the one or nucleic acid probes with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid of the sample with the one or nucleic acid with the one of the one of the one of the or nucleic acid 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Non-nucleotide probe for suppressing binding of detectable nucleic acid probes to undesired sequences, has aggregate nucleobase sequence homologous to randomly distributed repeat sequence of genomic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17
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ss; tumour suppressor; antitumour; cytostatic; tumour suppression; tumour regression; apoptosis; virus resistance; diagnosis; cellular degeneration.
                                                                                                                                                                                                                                                                                                                                ACC51496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               represents a
                                                                                                                               Human tumour suppressor sequence #263.
                                                                                                                                                                                                    27-JUN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 Similarity
17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATCTCGGCTCACTGCAA 983
                                                                                                                                                                                                                                                                                                                                   standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hyldig-Nielsen JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  non-nucleotide
                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NO 24; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 17; DB 1; L
Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Williams BF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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PT XXX PXX PF XXX PF XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New nucleic acid sequences associated with tumor suppression, regres apoptosis or virus resistance are useful to diagnose and treat viral disease, development of tumor cells and cell degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polypeptides having 80% identity to the polypeptide sequences. The invention is used to diagnose or treat viral disease or disease characterized by development of tumour cells or cellular degeneration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This sequence represents an isolated nucleic acid sequence associated with tumour suppression or regression, apoptosis or virus resistance. invention relates to these sequences or sequences having at least 80% identity to them, and polypeptides encoded by the sequences or minimum to them.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-JUN-2001; 2001FR-00008139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 101; 798pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-250498/25
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                                                 WPI; 2003-250498/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cellular degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ss; tumour suppressor; antitumour; cytostatic; tumour suppression; tumour regression; apoptosis; virus resistance; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACC54017;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACC54017 standard; DNA; 17
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                                                                                                    Tuijnder M,
                                                                                                                                                                                                            20-JUN-2001; 2001FR-00008139
                                                                                                                                                                                                                                                           20-JUN-2001; 2001FR-00008139
                                                                                                                                                                                                                                                                                                              27-DEC-2002.
                                                                                                                                                                                                                                                                                                                                                                  FR2826373-A1
                                                                                                                                                      (MOLE-) MOLECULAR ENGINES LAB
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17; Conserv
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SA
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1.4e+03
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New nucleic acid sequences associated with tumor suppression, regression

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RESULT 1183
ADL49972
ID ADL4997
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis restenosis; asthma; Crohn's disease; diabetes; obesit; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence represents an isolated nucleic acid sequence associated with tumour suppression or regression, apoptosis or virus resistance. invention relates to these sequences or sequences having at least 80% identity to them, and polypeptides encoded by the sequences or polypeptides having 80% identity to the polypeptide sequences. The invention is used to diagnose or treat viral disease or disease characterized by development of tumour cells or cellular degeneration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     apoptosis or virus resistance are useful to diagnose and treat viral disease, development of tumor cells and cell degeneration.
                                                                                                                                                                                                                                                          05-APR-2001; 2001US-00827395.
29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR) IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
                                                                                                         Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase opposes kinase PKR genes, for treating cancer and inflammatory disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADL49972 standard; RNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified.
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                                                                                                                                                                                                                              RIBOZYME PHARM INC.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        arthritis;
                                                                                                         kinase or
y disease.
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RESULT 1184
ADL50424
ID ADL5042
XX ADL5042
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Matches 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; central nervous system injury; CNS injury; spinal cord injury; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid ar restenosis; asthma; Crohn's disease; diabetes; obesity; arthumatoimnune disease; lupus; multiple sclerosis; transplant rejection; ischaemia; reperfusion; glomerulonephritis; se; allergy; asthma; allergic rhinitis; atopic dermatitis; human PK
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29-MAY-2001; 2001US-0294412P
28-AUG-2001; 2001US-0315315P
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13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                       Haeberli P,
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Pred. No.
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The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR), IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the

growth

enzymatic nucleic acid that down-regulates expression of neurite h inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase o in kinase PKR genes, for treating cancer and inflammatory disease

kinase or disease.

protein

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NO 3957;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                   antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis restenosis; asthma; Crohn's disease; diabettes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
                                                                                                                                                                                                                                                       05-APR-2001;
29-MAY-2001;
28-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human PKR substrate sequence #1846.
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                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
                                                                                                                                                                                            Blatt L,
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                                                                                                        protein kinase PKR
                                                                                                                                                                 WPI; 2003-058513/05
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                                             invention
                              down
               down regulate neurite growt
                                                                                                                     enzymatic nucleic
h inhibitor recepto
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13; Conserv
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                                                                                                                                                                                                                                                        2001US-00827395.
2001US-0294412P.
2001US-0315315P.
                                                                                                                                                                                                                                                                                                                    2002WO-US010512.
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comprises nucleic acids (e.g. antisense oliquilate the expression or inhibit the function of growth inhibitor, NOGO, prostaglandin D2 rece (IKK), or protein kinase PKR. The nucleic a
                                                                          ij
                                                                                                                     receptor,
                                                                          NO 4265;
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76.5%;
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                                                                                                                                                                                              Haeberli P,
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                                                                                                     acid that down-regulates expression of neurite
or, prostaglandin D2 receptor, IkappaB kinase ou
s, for treating cancer and inflammatory disease.
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Pred. No. 1.4e+03;
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tor (PTGDR)
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RESULT 1186
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                               antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IKappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; car melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthing restencesis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; ischaemia; reperfusion; glomerulonephritis; sepsiallergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
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                                                                                                                                            WPI; 2003-058513/05.
                                                                                                                                                                    Blatt L,
                                                                                                                                                                                                                       05-APR-2001; 2001US-00827395
29-MAY-2001; 2001US-0294412P
28-AUG-2001; 2001US-0315315P
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                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC.
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that down regulate the expression or for a neurite growth inhibitor, NOGO, IkappaB kinase (IKK), or protein kina

comprises nucleic acids (e.g. antisense oligonucleotides) unate the expression or inhibit the function of a receptor growth inhibitor, NOGO, prostaglandin D2 receptor (FTGDR), e (IKK), or protein kinase PKR. The nucleic acids of the

The invention

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2956;

317pp;

English

Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.

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RESULT 1187
ADL50218
ID ADL5021
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   The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor. NGGO, prostaglandin D2 receptor (PTGDR) IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                                                                                                                                                                                                                                     Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase opposed kinase PKR genes, for treating cancer and inflammatory disease
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nervous system (CNS)
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Pred. No.
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                                     a receptor
or (PTGDR)
                                                                                                                                                                                                                                                                                           kinase or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthriti restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor. NOGO, prostaglandin D2 receptor (PTGDR) IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the

Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.

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29-MAY-2001; 2001US-0294412P
28-AUG-2001; 2001US-0315315P
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                                       The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR)
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acids of

that down regulate for a neurite growt

growth inhibitor, NC (IKK), or protein

NOGO, prostaglandin D2 rec n kinase PKR. The nucleic A

din D2 receptor nucleic acids c

a receptor or (PTGDR)

(e.g. antisense oligonucleotides) inhibit the function of a receptor

The invention comprises nucleic acids (e.

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SEQ ID NO 2993; 317pp;

English.

Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.

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29-MAY-2001; 2001US-0294412P
28-AUG-2001; 2001US-0315315P
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             The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR)
                                                                                                                    Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR),

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                                                                                                                                                                                                Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
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Pred. No. 1.4e+03;
4; Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PYGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis restencesis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
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29-MAY-2001;
28-AUG-2001;
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                                                                                                                                                                                                            Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-APR-2002;
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      The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR) IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
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13; Conserv
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                                                                                                                                                                                                                                         05-APR-2001;
29-MAY-2001;
28-AUG-2001;
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or

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invention comprises nucleic acids (e.g. antisense oligonucleotides) toown regulate the expression or inhibit the function of a receptor

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Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disease, lupus, multiple sclerosis, transplant/graft rejection, ischaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The nucleic acids of the invention are also useful for down-regulating the expression of a target gene and as a diagnostic tool to examine genetic drifts and mutations within diseased cells or to detect the presence of target RNA in a cell. The present RNA sequence represents a human PKR
The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PFGDR) IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-APR-2001; 2001US-00827395.
29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-OCT-2002
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                                                                                                                                                                                                     enzymatic nucleic acid that down-regulates expression of neurite
h inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase o
in kinase PKR genes, for treating cancer and inflammatory disease
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29-MAY-2001; 2001US-0294412P
28-AUG-2001; 2001US-0315315P
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The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOZO, prostaglandin D2 receptor (PTGDR). IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the

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        The invention comprises nucleic acids (e.g. antisense oligibate down regulate the expression or inhibit the function for a neurite growth inhibitor, NOGO, prostaglandin D2 register that the function of the process of
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, prostaglandin D2 receptor, IkappaB kinase or
for treating cancer and inflammatory disease.
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Claim 59;

SEQ

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English.

Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.

WPI; 2003-058513/05.

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RESULT 1198
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29-MAY-2001; 2001US-0294412P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
   The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PRODR). IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
                                                                                                                                                                                                 Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antisense oligonucleotide; neurite growth inhibitor; NOGO; prosteglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cances melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthriti restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
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29-MAY-2001; 2001US-0294412P.
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The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor (PTGDR) for a neurite growth inhibitor, NGGO, prostaglandin D2 receptor (PTGDR) IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the

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                                      The invention comprises nucleic acids (e.g. antis that down regulate the expression or inhibit the for a neurite growth inhibitor, NOGO, prostagland IkappaB kinase (IKK), or protein kinase PKR. The
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                                                                                                                                                                       ID NO 3486; 317pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident;
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29-MAY-2001; 2001US-0294412P
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The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR)

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D2 receptor cleic acids o

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29-MAY-2001; 2001US-0294412P.
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The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR), IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
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                                                                                                                                                                                                                                                                                                                                                                                                                        RIBOZYME PHARM INC
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IkappaB kinase or
lammatory disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention are useful for treating: cerebrovascular accident, central nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma, lymphoma or glioma), inflammatory disease (e.g. rheumatoid arthritis, restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune disease, hupus, multiple sclerosis, transplant/graft rejection, ischaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The nucleic acids of the invention are also useful for down-regulating the expression of a target gene and as a diagnostic tool to examine genetic drifts and mutations within diseased cells or to detect the presence of target RNA in a cell. The present RNA sequence represents a human PKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-APR-2001;
29-MAY-2001;
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h inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or
in kinase PKR genes, for treating cancer and inflammatory disease.
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l Similarity 70.6%;
12; Conservative
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The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR) IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the

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                                                                                                                                                                                                                                                                                     05-APR-2001;
29-MAY-2001;
28-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PYGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 4 A;
                                                                                                                  Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                         03-APR-2002;
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The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR), IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
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; 2001US-0294412P.
; 2001US-0315315P.
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                                                                                    ID NO 4285; 317pp; English.
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RESULT 1206
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                                                                                                                                                                                                                                                                                                                                                                                                                   antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
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29-MAY-2001; 2001US-0294412P
28-AUG-2001; 2001US-0315315P
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prostaglandin D2 rese PKR. The nucleic

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(PIGDR)

invention comprises nucleic acids (e.g. antisense oligonucleotides) town regulate the expression or inhibit the function of a receptor

Novel enzymatic nucleic acid that down-regulates expression of neurit growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase protein kinase PKR genes, for treating cancer and inflammatory diseas

kinase or disease.

neurite

, prostaglandin for treating ca

or receptor, PKR genes,

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Claim 59;

SEQ ID NO

3487; 317pp;

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RESULT 1207
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ID ADL4997
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The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR) IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR) IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the

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2001US-0294412P.
2001US-0315315P.
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ches 0;
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RESULT 1209
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ID ADL4996
AC Unident
KW antisem
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
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29-MAY-2001;
28-AUG-2001;
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                                                                                                                                                  Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human PKR substrate sequence #1081.
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The invention comprises nucleic acids (e.g. antisense oligonucleotides) that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR) IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-APR-2002;
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13; Conserv
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2001US-0294412P.
2001US-0315315P.
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                                                                                                          ID NO 3500; 317pp; English.
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76.5%;
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Pred. No. 1.4e+03;
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RESULT 1210
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis restencesis; asthma; Crohn's disease; diabettes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
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29-MAY-2001; 2001US-0294412P
28-AUG-2001; 2001US-0315315P
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                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                    substrate; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antisense oligonucleotide; neurite growth inhibitor; NOGO;
prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human PKR substrate sequence #1867.
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                                                                                                                                                                                          Blatt L,
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inat down regulate for a neurite growt

growth inhibitor, NOGO, comprises nucleic acids (e.g. a ulate the expression or inhibit

The invention Claim 59;

SEQ ID NO 4286; 317pp;

English.

IkappaB

(IKK), or

protein

NOGO, prostaglandin D2 rekinase PKR. The nucleic

of.

antisense oligonucleotides)
t the function of a receptor
aglandin D2 receptor (PTGDR)

Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.

or

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RESULT 1211
ADK13213/c
ID ADK1321
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The present invention describes a method (M1) for aiding in the diagnosis of glioma. (M1) involves detecting an expression product of at least one gene (I) in a first brain tissue sample (T) suspected of being neoplastic, where (I) is chosen from any one of 255 genes (glioma endothelial markers (GEMs)) as given in specification, and comparing the expression of (I) in (T) with expression of (I) in a second normal brain tissue sample (R), where increased expression of (I) in (T) relative to (R), identifies (T) as likely to be neoplastic. Also described: (1) treating (M2) glioma involves contacting cells of the glioma with an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       glioma; brain tissue; neoplastic; glioma endothelial marker; GEM; anticancer; antiglioma; immune response; cytostatic; multi-drug sensitive glioma; human; long tag; ss.
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                                                                                                                                                                                                         genes, glioma endothelial markers, being neoplastic, and comparing th brain tissue sample.
                                                                                                                                                                                                                                         Diagnosing glioma by detecting expression product of any one of 255 genes, glioma endothelial markers, in brain tissue sample suspected of
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01-APR-2003; 2003US-0458978P.
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                                                                                                                                                                                                                                                                                                                             Wang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 4 A; 2 C; 7 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             endothelial marker (GEM) long tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                               S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.7%;
                                                                                                                                                                                                                                                                                                                               Cook BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score
Pred.
                                                                                                                                                                                                                         the expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                               Lattera
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17;
No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.4e+03;
                                                                                                                                                                                                                                                                                                                                 ŗ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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                                                                                                                                                                                                                         with expression in normal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO:391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>,,</u>
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RESULT 1212
ADK13231/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC antibody that specifically binds to a extracellular epitope; (2)
CC drug involves contacting a test compound with the cell which expresses
CC (I), monitoring an expression product of the at least one gene and
CC identifying test compound as a potential anticancer drug if it decreases
CC the expression of at least one gene; (3) identifying (M4) a test compound
CC as potential anticancer or antiglioma drug involves contacting a test
CC compound with the cell which expresses mRNA of at least one gene
CC identified by a tag as described above, monitoring mRNA of the gene, and
CC identifying the test compound as a potential anticancer drug if it
CC decreases the expression of at least one gene; and (4) inducing (M5) an
CC or (I). (I) have cytostatic activities, and can be used to trigger immune
CC useful for aiding in diagnosing glioma. (M2) is useful for treating multi
CC useful for aiding in diagnosing glioma. (M2) is useful for treating multi
CC cresponse to a glioma in a human. (M5) is useful for treating multi
CC adjioma surgically removed. The present sequence represents a human GEM
CC and tag oligonic lectide, which is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                   glioma; brain tissue; neoplastic; glioma endothelial anticancer; antiglioma; immune response; cytostatic; multi-drug sensitive glioma; human; long tag; ss.
                                                                     Diagnosing glioma by detecting expression product of any one of 255 genes, glioma endothelial markers, in brain tissue sample suspected of being neoplastic, and comparing the expression with expression in norm
                                                                                                                                                                                                                                      15-AUG-2002; 2002US-0403390P.
01-APR-2003; 2003US-0458978P.
                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human glioma endothelial marker (GEM)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADK13231;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             present invention.
                                                                                                                                                               Madden
                                                                                                                                                                                           (GENZ )
                                                                                                                                                                                                                                                                                                                                            WO2004016758-A2
                                                                                                                                                                                                                                                                                                                                                                                                             20-MAY-2004
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                                                          brain tissue sampie.
                                                                                                                                                                                                                                                                                 15-AUG-2003; 2003WO-US025614.
                                                                                                                                 2004-247973/23.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CCAAAGTGCTGGGATTA 403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCAAAGTGCTGGGATTA 1
                                                                                                                                                             Wang CJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                           HOPKINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5 C; 3
                                                                                                                                                             Cook BP,
                           114pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 17;
Pred. No.
                                                                                                                                                             Lattera J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                               Walter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 17;
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                                                                           normal
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Example 2;

SEQ

ID NO 409;

English

The present invention

describes a method

(M1)

for aiding

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diagnosis

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RESULT 1213
ADL82338/c
ID ADL8233
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ADL8233
AC ADL8233
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DT 20-MAY-
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Gene th
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US20031
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OB -JAN-
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PR 09-JAN-
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Best Local
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                                                                              Vector containing nucleic acid associated with breast cancer, useful treating, diagnosing and characterizing breast cancer, also related polypeptides and antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-JAN-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-SEP-2003
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                                                                                                                                                                                                                                                WPI; 2004-069003/07.
                                                                                                                                                                                                                                                                                                                                                                                                         (LYNX-) LYNX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l Similarity
17; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTACAGGCGCCCACCAC 756
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                         THERAPEUTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2003US-00339782.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     breast cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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309; 61pp; English

Sequence 17

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a composition which contains at least one vector (B) containing a nucleic acid (I) associated with breast cancer. The vector (B), also polypeptides (II) encoded by (I), are used for treatment of breast cancer. Arrays based on (I), (II), or their fragments, and (II) specific antibodies (Ab) are used to predict characteristics (e.g. invasiveness or stage) of breast cancer, and (I), or its fragments, are used to modulate characteristics of such cells; to identify breast cancer genes and to detect breast cancer (by detecting polymorphic nucleic acid or its products). The present sequence represents a human ER+ breast cancer differentially expressed sequence.
                          The invention relates to a novel method for identifying a subject at risl of breast cancer which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a risk of breast cancer, as well as therapeutic and prophylactic treatments that specifically target breast cancer, such as gene therapy. The current sequence is that of an Extend primer of the invention which was used to genotype single nucleotide polymorphisms within human glycoprotein VI (platelet) (GP6; GPIV;GPVI) DNA which is located at chromosomal position 19q13.4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         breast cancer; cytostatic; gene therapy; GP6; GPIV; GPVI; chromosome 19q13.4; ss; single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17
                                                                                                                                                                                                                                 Identifying a subject at risk of breast cancer by detecting the or absence of one or more nucleotide polymorphic variations, use diagnosing, preventing and/or treating breast cancer.
                                                                                                                                                                                                                                                                                                                                    Roth
                                                                                                                                                                                                     Example 3; Page 83; 286pp; English.
                                                                                                                                                                                                                                                                                                   WPI; 2004-441082/41
                                                                                                                                                                                                                                                                                                                                                                                                 25-NOV-2002;
24-JUL-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                25-NOV-2003; 2003WO-US037966
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Extend primer 60 used to genotype human glycoprotein VI polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-AUG-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADP08723
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                                                                                                                                                                                                                                                                                                                                                                  (SEQU-)
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17; Conserv
                                                                                                                                                                                                                                                                                                                                                                  SEQUENOM INC
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                                                                                                                                                                                                                                                                                                                                  Nelson MR,
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2003US-0490234P
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Pred. No.
                                                                                                                                                                                                                                                                                                                                      Kammerer
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hes 0;
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                                                                                                                                                                                                                                                                                                                                      SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human; platelet glycoprotein VI;
PCR; primer; SNP;
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RESULT 1215
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                                                    RESULT 1216
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Best Local S
Matches 17
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Best Local Similarity
Matches 17; Conser
                                                                                                                                                                                                                        The invention relates to a novel method for identifying a subject at risk of breast cancer which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a risk of breast cancer, as well as therapeutic and prophylactic treatments that specifically target breast cancer, such as gene therape. The current sequence is that of an Extend primer of the invention which was used to genotype single nucleotide polymorphisms within human glycoprotein VI (platelet) (GP6; GPIV;GPVI) DNA which is located at chromosomal position 19q13.4.
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  ADP08783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                                                                                                                                  Sequence 17
                                                                                                                                                                                                                                                                                                                                                                         Example 3; Page 82; 286pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                   diagnosing, preventing and/or treating breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                Identifying a subject at risk of breast cancer by detecting the presence or absence of one or more nucleotide polymorphic variations, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-441082/41.
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24-JUL-2003; 2003US-0490234P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           breast cancer;
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                           ADP08783 standard; DNA; 17
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17; Conserv
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                                                                                                                                               Conservative
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I; chromosome 19q13.4; ss; PCR; primer; SNP;
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; Pred. No. 1.40
0; Mismatches
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Pred. No.
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hes 0;
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RESULT 1217
ADP08787
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel method for identifying a subject at risk of breast cancer which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a risk of breast cancer,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            as well as therapeutic and prophylactic treatments that specifically target breast cancer, such as gene therapy. The current sequence is that of an Extend primer of the invention which was used to genotype single nucleotide polymorphisms within human glycoprotein VI (platelet) (GP6; GPIV;GPVI) DNA which is located at chromosomal position 19q13.4.
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                                                                                                                                                             breast cancer; cytostatic; gene therapy; GP6; GPIV; GPVI; chromosome 19q13.4; ss; single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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24-JUL-2003;
                                                                                                                                                                                                                                                                           Extend primer 124 used to genotype human glycoprotein VI polymorphism
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ilarity 100.0%;
Conservative (
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2003US-0490234P.
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Pred. No.
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PCR; primer; SNP;
                                                                                                                                                                                         human; platelet glycoprotein VI;
PCR; primer; SNP;
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WO2004047767-A2 Homo sapiens

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RESULT 1218
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Best Local :
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24-JUL-2003;
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24-JUL-2003;
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     Identifying a subject at risk of breast cancer by detecting the presence or absence of one or more nucleotide polymorphic variations, useful for
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                                                                                                                             Nelson
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2003US-0490234P.
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2003US-0490234P.
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MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               used to genotype human chromogranin B polymorphism.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 C;
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                                                                                                                                Braun
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7 G; 4 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 17;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kammerer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        breast cancer by detecting the presence tide polymorphic variations, useful for
                                                                                                                                   Kammerer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20pter-pl2; ss; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ,
KS
                                                                                                                                   MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer
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                                                                                                                                   Reneland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chromogranin B; CHGB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       a subject at risk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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diagnosing, preventing and/or treating breast cancer.
5
Page 102;
286pp; English
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The invention relates to a novel method for identifying a subject at risk of breast cancer which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a risk of breast cancer, as well as therapeutic and prophylactic treatments that specifically target breast cancer, such as gene therapy. The current sequence is that of an Extend primer of the invention which was used to genotype single nucleotide polymorphisms within human chromogranin B (CHGB; secretogranin 1,5CG1) DNA which is located at chromosomal position 20pter-p12.

Sequence 17 BP; 3 A; 6 C; 5 G; 3 T; 0 U; 0 Other;

Matches Query Match Local 17; Similarity Conservative 100.0%; 0 Score 17; Pred. Mismatches No. DB 1; Le 1.4e+03; Length 17 Indels 0 Gaps 0

맑 Ś 546 GCCTCCCAAGTAGCTGG 562 \_ GCCTCCCAAGTAGCTGG 17

RESULT 1219 AAH38113/c AAH38113 standard; DNA; 18 AAH38113;

specific upper PCR primer SEQ ID

14-AUG-2001

(first entry)

SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; lesch-Nyhan syndrome; muscular dystrophy; familial factorholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss. Single nucleotide polymorphism; SNP; single nucleotide primer extension;

Homo sapiens.

WO200129262-A2

26-APR-2001.

13-OCT-2000; 2000WO-US028436.

15-OCT-1999; 99US-0160096P

(ORCH-) ORCHID BIOSCIENCES INC

2001-290930/30 ŗ

HPI;

New genotyping oligonucleotide, absence or identity of single po e, useful for detecting the presence, polynucleotide polymorphism in a nucleic

Claim 1; Page 54; 83pp; English.

Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by representation and account of the summary automator resortion. single-nucleotide primer

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RESULT 1220
AAH91237/c
ID AAH9123
XX
AC AAH9123
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AC AAH9123
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CONTONION OF TOTONION
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HUMAN I
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HUMAN I
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Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis; single nucleotide polymorphism; SNP; chromosome 19p13; paternity test;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18
polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to the presence of genetic polymorphisms associated with inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human inflammatory bowel disease associated polymorphic site #312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                 (ELLI-)
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10-APR-2000; 2000US-0196046P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH91237;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH91237 standard;
                                                                                 The present invention describes a method for detecting the presence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-DEC-2000; 2000WO-US033632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-JUN-2001
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                                                                                                                                                                                        for the presence of polymorphisms sease, using a hybridization assay
                                                                                                                                                                                                                                                                                                                              Hudson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 4 A; 3 C; 7 G; 4 T; 0 U; 0 Other;
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                                                                                                                                      51; 463pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "SNP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polymorphism; SNI
3; forensic test;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
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                                                                                                                                                                                                                                                                                                                                 Lander
                                                                                                                                      English
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Pred. No.
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                                                                                                                                                                                              assay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.4e+03;
hes 0;
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RESULT 1221
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 7 A; 4 C; 2 G; 4 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                     The invention comprises a method for identifying a subject at risk of melanoma. The invention involves detecting the presence or absence of or more polymorphic variations associated with melanoma in the neuropi 1 (NRP1) or mannose receptor C type 2 (MRC2) genes. The method of the invention is useful for identifying subjects at risk and treating melanoma. The present DNA sequence represents an extension PCR primer that was used to detect single nucleotide polymorphisms within human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying a subject at risk of melanoma, useful for treating comprises detecting the presence or absence of one or more polyvariations associated with melanoma in a nucleic acid sample from the contract of the contract o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-NOV-2002; 2002US-0424475P.
23-JUL-2003; 2003US-0489703P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human neuropilin 1
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                                                                                                                                                                                                                                       Sequence 18
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17; Conserv
                                                                                                                             Similarity
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                   GGCTGGAGTGCAGTGGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 78; 176pp; English.
                                                                                                                                                                                                                                           BP;
                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                  present DNA sequence represents an extension to detect single nucleotide nolumnation
                                                                                                                                                                                                                                           2 A; 3 C; 9 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (NRP1)
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                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Braun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      extension
               663
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Pred. No. 1.4e
0; Mismatches
                                                                                                                             Score 17;
Pred. No.
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                                                                                              Mismatches
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                                                                                                                                                                 DB 1;
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                                                                                                                             1.4e+03;
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                                                                                                                                                                 Length 18;
                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sence of one neuropilin
                                                                                      Gaps
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RESULT 1222
ADO56522
ID ADO5652
RESULT 1223
ADD56536/c
ID ADD5653
XX
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X7
DT 12-AUG-
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DE Human c
XX
EW gene th
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                       melanoma comprising detecting the presence of absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a subject. Preventing melanoma in a subject comprises detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a subject; and administering a melanoma preventative to a subject in need thereof based upon the presence or absence of the one or more polymorphic variations in the nucleic acid sample. The preventative reduces ultraviolet (UV) light exposure to the subject. The methods, nucleic acids, proteins, and compositions are useful for treating melanoma. The present sequence represents a human cyclin-dependent kinase 10, CDK10, proximal SNP probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene therapy; human; ss; melanoma;
melanoma associated polymorphic variation; SNP;
single nucleotide polymorphism; cyclin-dependent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying a subject at risk of melanoma, useful for treating melanoma, comprises detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AD056522
                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a method of identifying a subject at melanoma comprising detecting the presence or absence of one of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example
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23-JUL-2003; 2003US-0489703P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-NOV-2003; 2003WO-US035879
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADO56522 standard; DNA; 18
 gene therapy;
                          Human cyclin-dependent kinase 10,
                                                      12-AUG-2004
                                                                                 ADO56536;
                                                                                                        AD056536
                                                                                                                                                                                                  392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cyclin-dependent kinase 10,
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                                                                                                                                                                                                                            l Similarity
17; Conserv
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••
                                                                                                                                                                                           GTGCTGGGATTACAGGC 408
                                                                                                                                                                                                                                                                                  18
                                                                                                          standard;
                                                                                                                                                                          GTGCTGGGATTACAGGC 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page
                                                                                                                                                                                                                                                                                BP; 3
                                                                                                                                                                                                                            Conservative
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                                                      (first entry)
    human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    84; 295pp;
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                                                                                                          DNA;
                                                                                                                                                                                                                                                                                  3 C; 7
    ss; melanoma;
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                                                                                                                                                                                                                                                                                  G; 4 T; 0 U; 1 Other;
                                                                                                                                                                                                                                          Score 17;
Pred. No.
                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CDK10 proximal
                           CDK10 proximal
                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                          1.4e+03;
                                                                                                                                                                                                                                                       Length 18
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                              SNP
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                             probe
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                              #61
                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            risk of
                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                              0;
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RESULT 1224 AAT65817/c

standard; DNA; 19

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25-MAR-2003 AAT65817; AAT65817

(revised)

entry)

17-JUN-1997

6

amplify repeat sequence marker Mfd10

repeat sequence; genetic marker; primer; amplification; se chain reaction; paternity; maternity; human; pedigree; sis; genetic disease; animal; plant; breeding; locus;

PCR; polymerase linkage analysis Polymorphism; Primer #2

analysis;

hybridisation;

chromosome;

gb

10-DEC-1996

Synthetic

뮍 8

17

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Query Match
Best Local
                                                                               Matches
                                                                                                                                                                                                                                       The invention relates to a method of identifying a subject at risk of melanoma comprising detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a subject. Preventing melanoma in a subject comprises detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a subject; and administering a melanoma preventative to a subject in need thereof based upon the presence or absence of the one or more polymorphic variations in the nucleic acid sample. The preventative reduces ultraviolet (UV) light exposure to the subject. The methods, nucleic acids, proceaus and compositions are useful for treating melanoma. The present sequence represents a human cyclin-dependent kinase 10, CDX10, proximal SNP probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying a subject at risk of melanoma, useful for treating melanoma comprises detecting the presence or absence of one or more polymorphic variations associated with melanoma in a nucleic acid sample from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-NOV-2002; 2002US-0424475P
23-JUL-2003; 2003US-0489703P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        melanoma аввосiated polymorphic variation; SNP;
Single nucleotide polymorphism; cyclin-dependent kinase 10; CDK10; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Roth
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                              Sequence 18 BP; 3 A; 5 C; 6 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-411721/38.
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                                                                                                      Local
851 GGCCTCCCAAAGTGCTG
                                                                               1 Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENOM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5; Page 85; 295pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nelson MR,
                                                                               Conservative
                                                                                                         100.0%;
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                          867
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                                                                               <u>;</u>
                                                                                                               Score 17;
Pred. No
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                                                                                    Mismatches
                                                                                                               No.
                                                                                                            1.4e+03;
                                                                                                                                      DB 1;
                                                                                                                                      Length 18;
                                                                                    Indels
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RESULT 1225
AAT49298
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   repeats, especially for use in e.g paternity or maternity testing, hu genetic analysis such as linkage analysis of genetic disease, commerce animal or plant breeding or pedigree analysis. Clones containing the repeat sequences were isolated by hybridisation of chromosome-specifi phage libraries with a synthetic poly(dC-dA). (dG-dT) probe. Over 100 repeat blocks were isolated. The primers AAT65798-T66047 were used to amplify the inserts from the isolated clones containing the repeat sequences. The primers AAT65816-7 were used to amplify the repeat sequence marker clone Mfd10 (AAT65712). (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to the isolation of polymorphic repeat sequences having the sequence (dC-dA)n. (dG-dT)n which can be used as genetic markers. Primers based on these sequences can be used to detect these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Detection of polymorphic genetic markers of the form (dC-dA)n(dG-dT)n using novel nucleic acid mols. as primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-APR-1994;
                                                                                                                                                                                                                                                                                                                                                        27-AUG-2003
27-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 4 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 7; Col 9-10; 186pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1997-042299/04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-APR-1989;
05-SEP-1991;
Detection of influenza virus endonuclease in a sample - by cleavage of RNA substrate to generate a primer for a labelled polymerase extension
                                             WPI; 1997-052364/05.
                                                                                                                                                              03-JUN-1996;
                                                                                                                                                                                                                                                   Alfalfa mosaic virus.
                                                                                                                                                                                                                                                                                                                           5' end fragment of Alfalfa Mosaic Virus 4.
                                                                                                                                                                                                                                                                                                                                                                                                    AAT49298;
                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT49298 standard; RNA; 19 BP
                                                                                                                                  07-JUN-1995;
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                                                                                                                                                                                                                         WO9640993-A1
                                                                                                                                                                                                                                                                                electrophoresis; substrate
                                                                                                                                                                                                                                                                                              Alfalfa Mosaic virus 4; influenza endonuclease; detection;
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                                                                                                      (MERI ) MERCK & CO INC.
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(first entry)
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91US-00754351.
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                                                                                                                                    95US-00487759
                                                                                                                                                              96WO-US008320
                                                                          Olsen DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.7%; Score 17;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                  cleavage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Lo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 19;
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RESULT 1226
AAT74905
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence was used as a substrate for influenza endonuclease in the method of the invention. The method allows detection of influenza endonuclease activity in a sample and comprises: (a) adding an influenza endonuclease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 6; Page 12; 28pp; English
                                                                                                                                                                                                                                                                                                                                                                                                             electrophoresis; substrate cleavage; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             5' end fragment of Alfalfa Mosaic Virus 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-AUG-2003
27-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 3 A; 1 C; 1 G; 0 T; 14 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence represents the 5' end of Alfalfa Mosaic virus 4 RNA.
             WPI; 1997-052365/05
                                                                                                                                                                                                                                                                                                                                                                            Alfalfa mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                              Alfalfa Mosaic virus 4; influenza endonuclease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT74905;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT74905 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              reaction.
                                            Cole JL,
                                                                                                             07-JUN-1995;
                                                                                                                                            03-JUN-1996;
                                                                                                                                                                                                                                                                           modified_base
                                                                                                                                                                                                                                                                                                                            modified_base.
                                                                                                                                                                            19-DEC-1996.
                                                                                                                                                                                                             WO9640994-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               601 TTTTTATTTTTAATTTT 617
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3; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 UUUUUAUUUUUAAUUUU 18
                                                                             MERCK & CO INC.
                                            Kuo LC, Olsen DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                            virus.
                                                                                                             95US-00487760.
                                                                                                                                              96WO-US008330.
                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                            /mod_base= Triphosphorylated-G
                                                                                                                                                                                                                                              mod_base= 2'-OMe-U
                                                                                                                                                                                                                                                               ′*tag= b
                                                                                                                                                                                                                                                                                                               *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 19;
1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                detection,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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RESULT 1227
AAT47271
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Best Local S
Matches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence was used in the method of the invention to the control activity in a sample. The method comprises: (a) adding an oligonucleotide substrate to a sample to generate an oligonucleotide product; (b) hybridising the oligonucleotide prod. with a DNA template which comprises a first segment complementary to the oligonucleotide and a 5' extension of at least one nucleotide attached to the 5' end of the DNA segment, such that a DNA:RNA hybrid or a DNA:DNA duplex is formed; (c) adding a DNA polymerase and labelled mononucleotides such that the DNA polymerase incorporates the mononucleotides to the 3' end of the oligonucleotide; incorporates the mononucleotides to the 3' end of the oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and (d) measuring the amt. of labelled hybrid prod. as a measure of the amt. of the enzyme activity in the sample. The method is used to assay for enzymes e.g. endonuclease, exonuclease or ribozymes, that act on substrates to generate single stranded oligonucleotide prods. by cleaving the substrate which then forms a primer for extension by a DNA polymerase on a template. It can be used to identify the position where the enzyme cleaves the substrate. The assay can also be used to screen for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detection of enzyme pref. endonuclease or ribozyme, in a sample - by cleavage of an RNA substrate to generate a primer for a labelled
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence represents the 5' end of Alfalfa Mosaic virus 4 RNA. This sequence was used in the method of the invention for detecting the enzyme activity in a sample. The method comprises: (a) adding an oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polymerase extension reaction.
                                                                                                                                                                                                                                                                                                                                                                                             Capped RNA molecule; mRNA maturation; translation initiation; influenza; endonuclease aptamer; RNase; therapy; inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Page 14; 34pp; English
07-JUN-1995;
                                                                                                                                                                                                         modified_base
                                                                                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                               Capped RNA influenza endonuclease substrate #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT47271 standard; RNA; 19
                                                                                                                                                           modified_base
                               03-JUN-1996;
                                                              19-DEC-1996
                                                                                              WO9640159-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inhibitors of these enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              601 TTTTTATTTTAATTTT
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3; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
95US-00480068
                                96WO-US008394
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 A;
                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                           /*tag=
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                                                                                                                               2'-deoxy-2'-fluoro-uridine
                                                                                                                                                                            2'-deoxy-2'-fluoro-uridine
                                                                                                                                                                                                                            2'-0-methyluridine
                                                                                                                                                                                                                                                                               triphosphorylated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G; 0 T; 14 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 17;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (Updated on 27-AUG-2003 to correct OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ç.
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AAT47276
AAT47276
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC the invention. The method of the invention is for producing capped RNA or CC RNA analogues. The method comprises reacting a RNA or analogue colligonucleotide with a phosphate addition agent to form a RNA or analogue colligonucleotide with a phosphate addition agent to form a RNA or analogue colligonucleotide with a phosphate which is then capped. The presence of the cap colligonucleotide with a phosphate addition of translation, and protects colligonucleotide to triphosphate, which is then capped. The presence of the cap is important for mRNA maturation, initiation of translation, and protects colligonucleotide with a maturation of the call. The capped RNA or canalogue is an influenza infection in an animal. The synthetic capped RNA are substrates for virally encoded endonuclease associated with influenza colliging. The short non-extendible (due to their length or because of the colligination of the 3' end of the oligo) RNA molecules are potent constitution of the short capped RNA by influenza endonuclease. They may be used to investigate viral and cellular mechanisms of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MERI ) MERCK & CO INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Capped RNA influenza endonuclease substrate #8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT47276;
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                              WPI; 1997-051868/05
                                                                                        Benseler F,
                                                                                                                                                                                                                                                             03-JUN-1996;
                                                                                                                                                                                                                                                                                                                    19-DEC-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               endonuclease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Capped RNA molecule;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-AUG-1997
                                                                                                                                            (MERI ) MERCK & CO INC
                                                                                                                                                                                                      07-JUN-1995;
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3; Conserve
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                                                                                        Cole JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               plecule; mRNA maturation; translation initiation; influenza;
aptamer; RNase; therapy; inhibitor; ss.
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                                                                                                                                                                                                      95US-00480068
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.7%;
17.6%;
                                                                                        Kuo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14;
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                                                                                        ť,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Olsen DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                        Olsen
                                                                                        DB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 19
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RESULT 1229
AAT47269
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Best Local S
Matches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA analogues. The method comprises reacting a RNA or analogue mono-, di- or triphosphate, which is then capped. The presence of the cap is important for mRNA maturation, initiation of translation, and protects the mRNA against various RNAses present in the cell. The capped RNA or analogue is an influenza endomuclease aptamer, useful for treating or preventing an influenza infection in an animal. The synthetic capped RNA are substrates for virally encoded endonuclease associated with influenza virus. The short non-extendible (due to their length or because of the modification of the 3' end of the oligo) RNA molecules are potent inhibitors of the cleavage of capped RNA by influenza endonuclease. They may be used to investigate viral and cellular mechanisms of
                                                                                                                                                                                                                                                                                                                                                                                            Capped RNA molecule; mRNA maturation; translation initiation; influenza; endonuclease aptamer; RNase; therapy; inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Production of capped RNA or analogues - useful as substrates for influenza virus associated virally encoded endonuclease.
                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                     Capped RNA influenza endonuclease substrate #3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT47269 standard; RNA; 19
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Production of capped RNA or analogues - useful as substrates for
                            WPI; 1997-051868/05
                                                                                                          07-JUN-1995;
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                                                      Benseler F,
                                                                                                                                     03-JUN-1996;
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                                                                                                                                                                                                                                            modified_base
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                                                                                  MERCK & CO INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                     Cole JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                           95US-00480068
                                                                                                                                      96WO-US008394
                                                                                                                                                                                                    /*tag= c
/mod_base= 2'-deoxyadenosine
                                                                                                                                                                                                                                                         /*tag= b
/mod_base=
                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                 /mod_base= triphosphorylated
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                                                        Kuo LC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14;
                                                                                                                                                                                                                                                            2'-0-methyluridine
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Pred. No.
                                                        Olsen DB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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SXXXXXX

07-JUN-1995;

95US-00480068 96WO-US008394

(MERI ) MERCK & CO INC

03-JUN-1996;

19-DEC-1996 WO9640159-A1 modified\_base

mod\_base= \*tag=

phosphorothioated

Ω,

phosphorothioated

\*tag=

a

modified\_base

'mod\_base= phosphorothioated

′\*tag= e mod\_base= modified\_base

/\*tag= b /mod\_base=

2'-O-methyluridine

mod\_base=

triphosphorylated

\*tag=

modified\_base

modified\_base

Location/Qualifiers

Synthetic

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RESULT 1230
AAT47279
ID AAT4727
XX
AC AAT4727
XX
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Best Local S
Matches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT47264-T47280 represent capped RNA molecules produced by the method of the invention. The method off the invention is for producing capped RNA or RNA analogues. The method comprises reacting a RNA or analogue oligonuclectide with a phosphate addition agent to form a RNA or analogue mono, di- or triphosphate, which is then capped. The presence of the cap is important for mRNA maturation, initiation of translation, and protects the mRNA against various RNases present in the cell. The capped RNA or analogue is an influenza endonuclease aptamer, useful for treating or preventing an influenza infection in an animal. The synthetic capped RNA are substrates for virally encoded endonuclease associated with influenza virus. The short non-extendible (due to their length or because of the modification of the cleavage of capped RNA by influenza endonuclease. They may be used to investigate viral and cellular mechanisms of
                                                                                                                                                                                                                                                                                                                                      Capped RNA molecule; mRNA maturation; translation initiation; endonuclease aptamer; RNase; therapy; inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 3 A; 1 C; 1 G; 0 T; 14 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 18; Page 13; 39pp; English.
                                                                                                                                                                                                                                                                                                                                                                                     Capped RNA influenza endonuclease substrate #11.
                                                                                                                                                                                                                                                                                                                                                                                                                 28-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT47279;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT47279 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              transcription/translation, or mRNA maturation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           influenza virus associated virally encoded endonuclease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        601
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
3; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             UUUUUAUUUUUAAUUUU 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTTTTATTTTAATTTT 617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 17; DB
Pred. No. 1.5e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                         influenza;
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RESULT 1231
AAT47277
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Best Local S
Matches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the invention. The method of the invention is for producing capped RNA or RNA analogues. The method comprises reacting a RNA or analogue oligonucleotide with a phosphate addition agent to form a RNA or analogue mono, di- or triphosphate, which is then capped. The presence of the cap is important for mRNA maturation, initiation of translation, and protects the mRNA against various RNAses present in the cell. The capped RNA or analogue is an influenza endonuclease aptamer, useful for treating or preventing an influenza infection in an animal. The synthetic capped RNA are substrates for virally encoded endonuclease associated with influenza virus. The short non-extendible (due to their length or because of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT47264-T47280 represent capped RNA molecules produced by the method of the invention. The method of the invention is for producing capped RNA of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Production of capped RNA or analogues - useful as substrates for influenza virus associated virally encoded endonuclease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Benseler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             virus. The short non-extendible (due to their length or because of the modification of the 3' end of the oligo) RNA molecules are potent inhibitors of the cleavage of capped RNA by influenza endonuclease. They may be used to investigate viral and cellular mechanisms of
                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                              Capped RNA molecule; mRNA maturation; translation initiation; influenza; endonuclease aptamer; RNase; therapy; inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                     Capped
                                                                                                                                                                                                                                                                                                                                                                                                                                    28-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT47277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT47277 standard; RNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transcription/translation, or mRNA maturation
Benseler F,
                                                                                                                                            WO9640159-A1
                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                         modified_base
                                                                                                                                                                                                                                                                                       modified_base
                                                       07-JUN-1995;
                                                                                  03-JUN-1996;
                                                                                                               19-DEC-1996.
                           (MERI ) MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                601 TTTTTATTTTTAATTTT 617
                                                                                                                                                                                                                                                                                                                                                                                                        RNA
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3; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                         influenza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 3 A; 1 C; 1 G; 0 T; 14 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cole JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry
Cole JL,
                                                         95US-00480068
                                                                                     96WO-US008394
                                                                                                                                                                                                                /*tag= b
/mod_base=
                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                   *tag=
                                                                                                                                                                                                                                                          mod_base=
                                                                                                                                                                       mod_base= 2'-0-methyluridine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.7%;
17.6%;
                                                                                                                                                                                                                                                                                                                                                                                                         endonuclease substrate #9.
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Kuo LC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
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                                                                                                                                                                                                                 2'-0-methyluridine
                                                                                                                                                                                                                                                           triphosphorylated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 17; DB 1; 1
Pred. No. 1.5e+03;
4; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Olsen DB
Olsen DB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RNA or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               production of capped RNA or analogues - useful as substrates
influenza virus associated virally encoded endonuclease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-051868/05
                                                                                                                                                                                                                                                                      Key
modified_base
                                                                                                                                                                                                                                                                                                                                        Capped RNA mo
                                                                                                                                                                                                                                                                                                                                                                             Capped RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 3 A; 1 C; 1 G; 0 T; 14 U; 0 Other;
                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                        28-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            transcription/translation, or mRNA maturation
                                                                                                                                                                                             misc_feature
                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT47273;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT47273 standard; RNA; 19
             Benseler F,
                                      (MERI ) MERCK & CO INC
                                                                07-JUN-1995;
                                                                                        03-JUN-1996;
                                                                                                                   19-DEC-1996.
                                                                                                                                          WO9640159-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    601 TTTTTATTTTAATTTT 617
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 3; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              UUUUUAUUUUUAAUUUU 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 15;
                                                                                                                                                                                                                                                                                                                                                     molecule; mRNA maturation; translation initiation; influenza;
                                                                                                                                                                                                                                                                                                                                                                               influenza endonuclease substrate #7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
             Cole JL,
                                                                                                                                                                                                                                                                                                                                         aptamer; RNase; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                95US-00480068
                                                                                         96WO-US008394
                                                                                                                                                                                                     /*tag= b
/mod_base=
                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                    /*tag=
/note=
                                                                                                                                                                                                                                      /*tag= a
/mod_base= triphosphorylated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        39pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.7%;
                                                                                                                                                            c
"biotin labelled for attachment
              Kuo LC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14;
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                                                                                                                                                                                                            2'-O-methyluridine
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Pred.
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              01sen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17;
              멺;
                                                                                                                                                                                                                                                                                                                                            inhibitor; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                       ç
                                                                                                                                                                       solid support"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
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RESULT 1233
AAT47264
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the invention. The method of the invention is for producing capped RNA or CRNA analogues. The method comprises reacting a RNA or analogue coligonucleotide with a phosphate addition agent to form a RNA or analogue mono, di- or triphosphate, which is then capped. The presence of the cap is important for mRNA maturation, initiation of translation, and protects che mRNA against various RNases present in the cell. The capped RNA or canalogue is an influenza infection in an animal. The synthetic capped RNA cresubstrates for virally encoded endonuclease associated with influenza cvirus. The short non-extendible (due to their length or because of the condification of the 3' end of the oligo) RNA molecules are potent completion/translation, or mRNA maturation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
Production of capped RNA or analogues - useful as substrates influenza virus associated virally encoded endonuclease.
                                         WPI; 1997-051868/05.
                                                                                                                          07-JUN-1995;
                                                                                                                                                    03-JUN-1996;
                                                                                                                                                                                                          W09640159-A1
                                                                                                                                                                                                                                                              modified_base
                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                         Capped RNA molecule; mRNA maturation; translation initiation; influenza; endonuclease aptamer; RNase; therapy; inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                  5' fragment of alfalfa mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                              27-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT47264 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT47264-T47280 represent capped RNA molecules produced by the method of
the invention. The method of the invention is for producing capped RNA or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 18; Page 14; 39pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Production
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1997-051868/05
                                                                                                                                                                                                                                                                                                      modified_base
                                                                                              (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          influenza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.7%;
Local Similarity 17.6%;
hes 3; Conservative
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                                                                   '13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of capped RNA or analogues - useful as substrates virus associated virally encoded endonuclease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 3 A; 1 C; 1 G; 0 T; 14 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                   Cole JL,
                                                                                                                         95US-00480068
                                                                                                                                                    96WO-US008394
                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                   'mod_base= 2'-O-methyluridine
                                                                                                                                                                                                                                                  /*tag= b
                                                                                                                                                                                                                                                                           mod_base= triphosphorylated
                                                                                                                                                                                                                                                                                          *tag=
                                                                    Kuo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВP
                                                                   ŗ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 17;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                    Olsen DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.5e+03;
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AAT47272
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 3
                                                                                                                                                                                                                                                                                                                                             Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Capped RNA
                            07-JUN-1995;
                                                          03-JUN-1996;
                                                                                                                                                                      modified_base
                                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                        Capped RNA molecule; mRNA maturation; endonuclease aptamer; RNase; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-AUG-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT47272 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 3 A; 1 C; 1 G; 0 T; 14 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 18; Page 12; 39pp; English
                                                                                     19-DEC-1996
                                                                                                                  WO9640159-A1
                                                                                                                                                                                                                                                                                                    modified_base
(MERI ) MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           601 TTTTTATTTTAATTTT 617
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Pred. No. 1.5e+03;
.4; Mismatches 0;
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Benseler F, Cole JL,

Kuo LC,

Olsen DB;

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RESULT 1235
AAT47278
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Matches 3; Conserv
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               Benseler F,
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             Cole JL,
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17.6%;
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Pred. No. 1.5e+03;
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                DB;
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RESULT 1236
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endonuclease aptamer; RNase; therapy;
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Production of capped RNA or analogues - useful as substrates influenza virus associated virally encoded endonuclease.
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3; Conserv
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17.6%; Pred. No.
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4; Mismatches
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RESULT 1237
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            Production of capped RNA or analogues - useful as substrates influenza virus associated virally encoded endonuclease.
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RESULT 1238
AAT63215/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT47264-T47280 represent capped RNA molecules produced by the method of the invention. The method of the invention is for producing capped RNA or RNA analogues. The method comprises reacting a RNA or analogue coligonuclectide with a phosphate addition agent to form a RNA or analogue mono, di- or triphosphate, which is then capped. The presence of the cap is important for mRNA maturation, initiation of translation, and protects the mRNA against various RNases present in the cell. The capped RNA or analogue is an influenza endonuclease aptamer, useful for treating or preventing an influenza endonuclease aptamer, useful for treating or preventing an influenza infection in an animal. The synthetic capped RNA are substrates for virally encoded endonuclease associated with influenza virus. The short non-extendible (due to their length or because of the inhibitors of the cleavage of capped RNA molecules are potent inhibitors of the cleavage of capped RNA by influenza endonuclease. They may be used to investigate viral and cellular mechanisms of
            Inter-Alu PCR was performed on YACs 905C2 and 763B11. Unpurified YAC DNA was amplified with generate primers Alu 5' (AAT63214) and Alu 3' (AAT63215). Genetic linkage strategies have placed a gene causing early onset Alzheimer's disease (AD) on the long arm of chromosome 14 between D14S289 and D14S61. The gene, S182 (see also AAT63207), was localised to a 100 kb region between D14S77 and D14S668E (see also AAT63216-22). A
                                                                                                                                                              New mutants of the S182 gene associated with familial Alzheimer's disease - and related protein and transgenic animals, useful as models for screening and assessing potential drugs.
                                                                                                                                                                                                                                                                                                                                                        18-JUL-1995;
02-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                             06-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               S182 gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT63215 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
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                                                                                                                               Example 2; Page 11; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                         26-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Primer Alu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT63215;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       UUUUUAUUUUUAAUUUU
                                                                                                                                                                                                                                                                     Hardy JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              familial Alzheimer's
chain reaction; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                      WASHINGTON SCHOOL SOUTH FLORIDA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or mRNA maturation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 17; DB 1;
Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                          MED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease; diagnosis; transgenic animal;
primer; artificial chromosome; PAC; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for PAC isolation
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number of novel

mutations

in the

S182 gene

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RESULT 1239
AAA35946/c
ID AAA3594
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Best Local Similarity
Matches 14; Conserv
                                                                    Query Match
Best Local
                                                        Matches
                                                                                                                                      A method has been developed for detecting the presence or absence of a single nucleotide polymorphism (SNP) allele in a genomic sample. The method comprises preparing a reduced complexity genome (RCG) from the genomic sample and analysing the RCG for the presence or absence of a SNP allele. The method can be used to characterise a tumour, to generate a genomic pattern for an individual genome or to generate a genomic classification code for a genome. The method can be used to assess whether a subject is at risk for developing a disease or to identify a set of SNP alleles associated with a disease. The method can also be used to perform linkage analysis. AAA35944 to AAA35947 represent sequences used in the exemplification of the present invention. AAA35948 to AAA36632 represent nucleotide sequences containing SNPs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               families multiply affected by early onset
                                                                                                                                                                                                                                                                                                                                                                               Detection of single nucleotide polymorphisms in genomes by preparation and analysis of reduced complexity genomes, useful for genotyping,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-APR-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Alu PCR primer 8C used for identifying
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA35946;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA35946 standard;
                                                                                                                                                                                                                                                                                                                                   Example 1; Page 75; 111pp;
                                                                                                                                                                                                                                                                                                                                                             and analysis of reduced complexity genomes, useful fingerprinting and determining allele frequency of
                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-293181/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MASI ) MASSACHUSETTS INST TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            allele
                                                                                                               Sequence 19 BP; 4 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               651 GGAGTGCAGTGGCGCAATC 669
                           967
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single nucleotide polymorphism; SNP; genotyping; DNA analysis; specific oligonucleotide; ASO; reduced complexity genome; RCG;
18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            characterisation; hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      JE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      specific oligonucleotide; Asu; remuser in specific oligonucleotide; Asu; remuser in classification; DNA fingerprinting; classification; BNA fingerprinting;
                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGAGTGCARTGGYRYRATC 1
                            ATCTCGGCTCACTGCAA 983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 3 A; 6 C; 2 G; 3 T; 0 U; 5 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jordan
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                                                        Conservative
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73.7%;
                                                                    1.7%;
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                                                                                                                                                                                                                                                                                                                                     English.
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                                                                    Score 17;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             No. 1.5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Charest
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                     DB 1;
                                                                       1.5e+03;
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                                                                                   Length 19;
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                                                           Indels
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                                                                              ABK93751
                                                                                           RESULT 1241
                                                                                                                                                                        Matches
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Best Local
                                                                                                                                                                                                                                                       The present invention describes a method for detecting the presence of polymorphisms associated with inflammatory bowel diseases such as polymorphisms associated with inflammatory bowel diseases. The methods can be used to detect the presence of genetic polymorphisms associated with inflammatory bowel disease and correlating their occurrence with disease states. They may be used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; inflammatory bowel disease; Crohn's disease; ulcerative cosingle nucleotide polymorphism; SNP; chromosome 19p13; paternity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH91329;
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                                                                   ABK93751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-DEC-1999; 99US-0170257P.
10-APR-2000; 2000US-0196046P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                       Testing for the presence of polymorphisms associated with bowel disease, using a hybridization assay.
                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-367874/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-JUN-2001.
Human inhibitor of apoptosis,
                        26-AUG-2002
                                                                                                                                                                                                                     Sequence 19
                                                                                                                                                                                                                                            polymorphic site described in the exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200142511-A2
                                              ABK93751;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ELLI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (WHED )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inflammatory bowel disease associated polymorphic site
                                                                                                                           19
                                                                                                                                                                       l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WHITEHEAD INST BIOMEDICAL RES. ELLIPSIS BIOTHERAPEUTICS CORP.
                                                                                                                                                                                                                                                                                                                                                   Page 55; 463pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Hudson
                                                                     standard;
                                                                                                                             TTTTNGAGACAGAGTCT
                                                                                                                                            TTTTTTGAGACAGAGTCT
                                                                                                                                                                                                                      BP; 8 A; 4 C; 2 G; 4 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000WO-US033632
                                                                                                                                                                         Conservative
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                        (first entry)
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/note= "SNP,
position"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polymorphism; SNF
3; forensic test;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                    1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Lander ES,
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                                                                                                                                                    631
                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               optional insertion or deletion
                                                                                                                                                                                    Score 17; DB 1;
Pred. No. 1.5e+03;
  HIAP1,
                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                    Rioux J,
                                                                                                                                                                                   No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene therapy;
  antisense oligonucleotide #2.
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                                                                                                                                                                                               Length 19;
                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                of.
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                                                                                                                                                                                                                                                                                                                                                                                        inflammatory
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RESULT 1242
ABZ75622/c
ID ABZ7562
XX ABZ7562
XX ABZ7562
XC ABZ7562
XC ABZ7562
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XX ABZ7562
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                                                                                                                                                                                                                                                                      吊
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an inhibitor of apoptosis (IAP) antisense nucleic acid (I) that inhibits IAP biological activity, regardless of length of the antisense nucleic acid, the IAP proteins may be mouse or human XIAP, HIAP1 or HIAP2. Also included are a pharmaceutical composition comprising a mammalian IAP antisense molecule and a method of enhancing apoptosis in a cell, comprising administering a negative regulator of the IAP anti-apoptotic pathway to the cell. The IAP antisense inhibitor is useful for enhancing apoptosis in a cell in a matisense inhibitor is useful for enhancing apoptosis in a cell in a matisense inhibitor is useful for enhancing apoptosis in a cell in a mammal diagnosed with a proliferative disease. The method is useful for treating a patient diagnosed with a proliferative disease in the cancer. The IAP antisense molecule is useful to treat, ameliorate, improve, adenocarcinoma, lymphoma, pancreatic cancer,) and also in diseases or conditions where apoptosis is involved or implicated (e.g. embryonic devalues).
                                                                                                                                                                                                                                                                                                                                             Best Local Similarity Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cytostatic; cancer; ovarian cancer; adenocarcinoma; lymphoma; IAP pancreatic cancer; embryonic development; viral pathogenesis; autoimmune disorder; neurodegenerative disease; multiple sclerosi lupus erythematosus; herpes virus infection; pox virus infection; adenovirus infection; proliferative disease.
Aneuploidy; chromosome; multiplex assay; polymerase chain reaction; short tandem repeat; STR; turner syndrome; cystic fibrosis; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         development, viral pathogenesis, autoimmune disorders, neurodegenerative diseases, multiple sclerosis, lupus erythematosus and infection by herpe diseases, pox virus and adenovirus). The present sequence is an IAP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 9; Page 36; 135pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel antisense inhibitor of apoptosis nucleic acid useful for enhancing apoptosis in a cell, for treating cancer and other proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-APR-2002
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                                                         STR marker 21-32S specific PCR primer 32S forward.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Korneluk RG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-SEP-2000; 2000US-00672717.
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                                                                                                15-MAY-2003
                                                                                                                                                                       ABZ75622 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (AEGE-)
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                                                                                                                                                                                                                                                                                                         CTCCTGCCTCAGCCTCC 551
                                                                                                                                                                                                                                                                    CTCCTGCCTCAGCCTCC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                               molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 5 A; 3 C; 10 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Baird S,
                                                                                                                                                                         ВÞ
                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                   Score 17;
Pred. No.
                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Holcik M,
                                                                                                                                                                                                                                                                                                                                                                 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                               ٥,
                                                                                                                                                                                                                                                                                                                                                                                  Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Young
                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            lymphoma; IAP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sclerosis;
                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 by herpes
                                                                                                                                                                                                                                                                                                                                               Gaps
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to detecting aneuploidy of a chromosome and involves using a multiplex polymerase chain reaction assay having chromosome-specific short tandem repeat (STR) markers. The STR marker 21-328 (informal designation) is useful as a marker for the diagnosis of aneuploidy of a chromosome, particularly trisomy 21, 13, 18 or X, or Turner Syndrome. The STR marker Y-408 (informal designation) is useful as a marker for the diagnosis of the sex of an individual. Marker CF508 is useful for detecting the presence or absence of a genetic disease, particularly cystic fibrosis. Sequences ABZ75621-22 represent PCR primers specific for the STR marker 21-32S
                                                                                                                                                                 Synthetic
                                                                                                                                                                                        Capped RNA molecule; mRNA maturation;
endonuclease aptamer; RNase; therapy;
                                                                                                                                                                                                                          5' fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting aneuploidy of a chromosome in a fetus by using a multiplex polymerase chain reaction assay comprising chromosome-specific short
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-707013/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-FEB-2001; 2001GB-00004690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-FEB-2002; 2002WO-GB000839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-SEP-2002
             WO9640159-A1
                                                        modified_base
                                                                                          modified_base
                                                                                                                                                                                                                                                   27-AUG-1997
                                                                                                                                                                                                                                                                         AAT47265;
                                                                                                                                                                                                                                                                                                AAT47265 standard; RNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 16; 30pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tandem repeat markers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Levett LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200268685-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                           modified_base
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                                                                                                                                                                                                                                                                                                                                                                                638
                                                                                                                                                                                                                                                                                                                                                          17
                                                                                                                                                                                                                                                                                                                                                                                                      l Similarity
17; Conserv
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                                                                                                                                                                                                                                                                                                                                                          TGTCACCCAGGCTGGAG
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                                                                                                                                                                                                                          #2 of alfalfa mosaic virus
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                                                                                                                                                                                                                                                  (first entry)
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                                                                    mod_base=
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                                                                                                                                                                                                                                                                                                                                                                                                     k; Score 17; DB
k; Pred. No. 1.5
0; Mismatches
                                   2'-O-methyluridine
                                                                                                       7-methylguanosine
                                                                     triphosphorylated
                                                                                                                                                                                                                                                                                                                                                                                                                                                    G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                         inhibitor;
                                                                                                                                                                                                      translation initiation;
                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                            Length 19;
                                                                                                                                                                                           88
                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                      influenza;
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RESULT 1244
AAZ37711/c
ID AAZ3771
XX AAZ3771
XX AAZ3771
XX Human m
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT47264-T47280 represent capped RNA molecules produced by the method of CC the invention. The method of the invention is for producing capped RNA or CC RNA analogues. The method comprises reacting a RNA or analogue CC oligonuclectide with a phosphate addition agent to form a RNA or analogue CC mono- di- or triphosphate, which is then capped. The presence of the cap CC is important for mRNA maturation, initiation of translation, and protects CC the mRNA against various RNAses present in the cell. The capped RNA or CC analogue is an influenza endonuclease aptamer, useful for treating or CC preventing an influenza infection in an animal. The synthetic capped RNA CC are substrates for virally encoded endonuclease associated with influenza CC virus. The short non-extendible (due to their length or because of the CC modification of the 3' end of the oligo) RNA molecules are potent CC inhibitors of the clavage of capped RNA by influenza endonuclease. They cap be used to investigate viral and cellular mechanisms of transcription/translation, or mRNA maturation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Production of capped RNA or analogues - useful as substrates influenza virus associated virally encoded endonuclease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1997-051868/05
                                                                                                                                                                                                                                                                                                                                                                                                   Human mdm2 gene; proliferation; tumour; phosphorothioate; p53; antisense; modulation; oligomucleotide; expression; inhibition; hyperproliferation; blood cancer; brain cancer; breast cancer; burg cancer; soft tissue cancer; psoriasis; fibrosis; atherosclus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Benseler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human mdm2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ37711;
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                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                               WO9949065-A1.
                                 (ISIS-)
                                                                                      26-MAR-1998;
                                                                                                                                        26-MAR-1999;
                                                                                                                                                                                             30-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                               restenosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
nes 3; Conserv
                                                                                                                                                                                                                                                                                            sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      phosphorothioate oligodeoxynucleotide #241.
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                                                                                      98US-00048810.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 A;
                                                                                                                                        99WO-US006702.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             39pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 C; 2 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 17; DB 1;
Pred. No. 1.5e+03;
4; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Olsen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         p53; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>,</u>
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RESULT 1245
AAA96372/c
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           novel nucleotide antisense compounds, targetted to the 5' untranslated, translation termination codon, or 3' untranslated region of a nucleic acid encoding human mdm2, that modulates expression of human mdm2. The oligonucleotides mediate their effect by antisense inhibition of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Miraglia LJ, Nero P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hyperproliferative gene expression. The antisense compound is used to treat an animal having a disease or condition associated with mdm2, particularly a hyperproliferative condition, more particularly cancer, especially of the blood, brain, breast, lung or soft tissue, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ37471, AAZ37472, AAZ37739, AAZ37740 and AAZ37741 are used in the exemplification of the present invention. The present function of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ37471, AAZ37472, AAZ37739, AAZ37740 and AAZ37741 are used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compounds used to treat
                                                                                                                                                                                                                                                                                                                                            ICOS gene; CTLA4 gene; costimulatory receptor gene locus; CGRL; lupus; insulin-dependent diabetes mellitus; IDDN; Addison's disease; leprosy; Graves disease; autoimmune hypothyroidism; myasthenia gravis; thymoma; thyroiditis; postpartum thyroiditis; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                       24-MAR-2000; 2000WO-US007938
                                                                                                                                                                                                                                                                              WO200056856-A2
                                                                                                                                                                                                                                                                                                                                                                                                Autoimmune disease; polymorphic microsatellite repeat; PMR;
                                                                                                                                                                                                                                                                                                                                                                                                                            Primer used to amplify a sara3/4 polymorphic microsatellite repeat.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA96372;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA96372 standard; DNA;
                                                                                                                                                                                                                                                   28-SEP-2000
                                                                                                                                                                                                                                                                                                                                   Hashimoto's disease; coeliac disease; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1999-610754/52
                                                                                                                                                                                                                                                                                                        sapiens.
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17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              fibrosis, atherosclerosis or restenosis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 7 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 17;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    eg.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hyperproliferative
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                                                                                                                                                                                                                                                                                                                                                                                                       CD28 gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                         lupus;
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PCR primers AAA96371-72 were used to amplify polymorphic microsatellite repeat (PMR) sequences from the human costimulatory receptor gene locus

Claim 18;

Page

147;

160pp;

English

Determining predisposition of humans to develoninvolves detecting polymorphic microsatellite

develop autoimmune disease ellite repeat sequence within

costimulatory

receptor gene locus.

WPI: Ling

2000-628257/60.

25-MAR-1999;

99US-0126215P

(GEMY)

GENETICS INST INC

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Gray

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RESULT 1246
AAC59889/c
ID AAC5988
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Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-MAR-1999;
17-AUG-1999;
01-OCT-1999;
   This invention relates to 59 human secreted proteins and the nucleotide sequences encoding them. Sequences AAC59788-C59846 and AAB34687-B34745
                                                                                                                                                           Novel proteins and polypeptides useful for the treatment of e.g multiple sclerosis, systemic lupus erythmatosus, rheumatoid arthritis, cancer, Alzheimer's disease, Parkinson's disease, stroke, anemia and ulcers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Secreted protein; human; autoimmune disorder; multiple sclerosis; ulcer; systemic lupus erythematosus; rheumatoid arthritis; anaemia; stroke; haematopoiesis regulation; tissue regrowth; wound healing; haemophilia; hlzheimer s disease; Parkinson's disease; Shy-drager syndrome; cancer; contraceptive; infection; growth inhibition; hyperproliferative disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide probe for human DNA clone vq9 1.
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                                                                                            Disclosure; Page 472; 493pp; English
                                                                                                                                                                                                                                                                                                                                                        Valenzuela D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                       ALPHAGENE INC.
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99US-0149639P.

99US-0157247P.

99US-0167824P.

2000US-0182711P.
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                                                                                                                                                                                                                                                                                                                                                        Yuan O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0124808P
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                                                                                                                                                                                                                                                                                                                                                        Rapiejko
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RESULT 1247
AAK94972
ID AAK9497
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                  08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 4 A; 9 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                07-JUL-2000; 2000EP-00114089.
                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                       Human; full length
                                                                                                                                                                                                                                                                                                                                                                                                                Human cDNA clone-specific primer, SEQ ID NO: 4217.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAK94972;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        867 GGGATTACAGGCGTGAG 883
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                                  Nishikawa T,
su A, Sugiyama
                                                                                              HELIX RES INST.
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                                                                                                                                  2000JP-00118774.
2000JP-00183765.
                                                                                                                                                                         99JP-00194486.
                                                                                                                                                                                                                                                                                                                                                                         cDNA; cDNA synthesis; oligo-capping; PCR
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                                  Isogai T,
na T, Nagai
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Pred. No. 1.5e+03
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                                    Hayashi K,
K, Kojima
                                                     Ishii S,
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                                      Kawai Y;
T, Koga
                                                                                                                                                                                                                                                                                                                                                                            primer;
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RESULT 1248
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful length clones were obtained by construction of the protein encoded by the cDNA. The full libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA easily without any special methods. The present sequence is a primer used to amplify a human cDNA clone provided in the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           830\ \text{Primers} useful for synthesizing full length cDNA clones and their in genetic manipulation.
                      The present invention relates to an antisense compound 8-30 nucleobases in length targeted to nucleobases 1-308 of the 5' untranslated region, 1776-1806 of the translation termination codon region or 1818-2370 of the 3' untranslated region of a nucleic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro. The hyperproliferative disorder includes cancer or psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 18; Page 127; 1380pp +
                                                                                                                                                                        Novel antisense compound 8-30 acid molecule encoding human
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                                                                                                                                 Example
                                                                                                                                                            of human
                                                                                                                                                                                                                WPI; 2001-190948/19
                                                                                                                                                                                                                                          Miraglia
                                                                                                                                                                                                                                                                                                26-MAR-1998;
                                                                                                                                                                                                                                                                                                                          26-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense; mdm2; hyperproliferation; cancer; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human
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17; Conserv
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                                                                                                                                                          mdm-2 and reducing hyperproliferation
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                                                                                                                               Col 31; 77pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                            phosphorothicate oligonucleotide #239.
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                                                                                                                                                                                                                                                                     PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                          Nero P,
                                                                                                                                                                                                                                                                                                98US-00048810
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                                                                                                                                                                                                                                          Graham
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                                                                                                                                                            8-30 nucleobases in length targeted to a nucle
uman mdm-2 useful for modulating the expression
ing hyperproliferation of human cells.
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Pred. No.
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5. 1.5e+03;
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Query Match

Sequence

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Score 17;

<u>ب</u> Other;

Length 20,

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XX Human m
XX Human;
XW Human;
XW Atheros
XX Homo sa
XX FF ROJ
FT modific
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XX (MIRA/)
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The present invention relates to antisense compounds, 8-30 nucleobases in CC length targeted to the 5' untranslated region, translation termination CC codon region, 3' untranslated region, coding region or translation start CC site of a nucleic acid encoding human mdm2 where the antisense compound CC modulates the expression of human mdm2. The antisense oligonucleotides of the invention are useful for encoding human mdm2 and for inhibiting the CC expression of human mdm2. They may be used for treating an animal having CC adisease or condition associated with amplification of mdm2 gene or CC (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis, fibrosis, atheroselerosis or restenosis, tumours, colorectal carcinoma CC and chronic myelogenous leukemia. The antisense compound may be administered with a chemotherapeutic agent to overcome drug resistance. CC The antisense compound reduces hyperproliferation of human cells. The method, which involves the use of the antisense compound, is also useful CC method, which involves the use of the antisense compound, is also useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; mdm2; hyperproliferative disorder; cancer; ps
atherosclerosis; tumour; cytostatic; anti psoriatic;
anti arteriosclerotic; vasotropic; antisense; phosph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    An antisense compound, useful for nucleobases targeted a region (e. of a nucleic acid encoding human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAS29480 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example
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         This invention relates to the CDNA and protein sequences of novel proteins HYPLIP1 or FCHII and to sequence variations within these genes that have been shown to be associated with lipid disorders.

Chat have been shown to be associated with lipid disorders.

Coligonucleotide probes that hybridise to the CDNA sequence are useful for analysing the expression of FCHII by detecting the expression of the mRNA transformed with the CDNA of the invention is useful for producing the protein by recombinant means.

Charlest in the sample. A host cell transformed with the CDNA of the invention are useful for producing the protein by recombinant means.

Charlest in the sample and the sequence of the invention are useful for treating or preventing a lipid disorder associated with expression of FCHII such as familial combined hyperlipidaemia, coronary are useful for treating or preventing a lipid disorder associated with hyperapobetalipoproteinaemia, hypertriglyceridaemia, familial dyslipidaemic hypertension, syndrome X, obesity, insulin resistance and hypercholesterolaemia. The cDNA sequence is useful in the diagnosis or prognosis of predisposition to lipid disorders and cancers, and also to identify a molecule which enhances or decreases the HYPLIP1 or FCHII activity. The prosent sequence represents an oligonucleotide primer activity. The prosent sequence represents an intention The mouse HypLIP1 locus of the intention.
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ARESULT 1251
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                                                    The present invention describes a method of arraying genome clones. The CC method comprises: (a) clones of the genomic libraries contained in CC multiwell plates numbered for discrimination are mixed in each of the CC multiwell plates; (b) a primer designed based on the chromosome marker (c) a signal corresponding to the maxture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant cC mplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order cc of the maximum in the specified discrimination Nos. to array the multiwell cc plates; (e) the clones in the multiwell plates of the specified cd discrimination Nos. are mixed respectively in each wells of longitudinal and lateral directions; (f) the mixed clones are cultured and the cc resultant cultures are amplified by using the above primer; (g) signals are detected from the amplified by using the above primer; (g) signals are detected from the detected result; and (i) the clones are mixed respectively. The clones in the multiwell plates are specified from the detected result; and (i) the clones are constituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABL42957 to ABL45322 represent CC specifically claimed for use in the present invention
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Query Match
Best Local Similarity
Matches 17; Conserv

Conservative

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Mismatches

Indels

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Gaps

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100.0%;

Score 17; Pred. No.

DB 1;

Length 20

1.5e+03

20

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5 C; 5 G; 4 T; 0 U; 0 Other;

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RESULT 1252
ABK71108
RESULT 1253
AAD52338
ID AAD5233
XX AAD5233
XX AAD5233
XX O2-MAY-
XX O2-MAY-
XX Human I
XX Antisen
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                                                                                                                                                    Matches
                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                     The invention relates to an isolated polynucleotide comprising a sequence variation of a mouse HYPLIP1 cDNA or a human FCHL1 (familial combined hyperlipidaemia) gene. The FCHL1 polynucleotide, the FCHL1 polypeptide or antibody immunoreactive to the FCHL1 polypeptide are useful for treating or preventing cancer associated with expression of FCHL1, as well as for treating lipid disorder. The mouse HYPLIP1 cDNA or human FCHL1 gene are also useful for diagnosing or prognosing a predisposition to lipid disorder and cancer. ABK70902-ABK71303 represent mouse HYPLIP1, human FCHL1 coding sequences and PCR primers of the invention
                                                                                                                                                                                                                                                                                                              New mouse HYPLIP1 and human FCHL1 (familial genes and their sequence variations, useful preventing lipid disorders and cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; mouse; HYPLIP1; FCHL1; familial lipid disorder; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABK71108 standard; DNA; 20
                                                                                                                                                                                      Sequence 20 BP; 5 A;
                                                                                                                                                                                                                                                                                             Claim 11; Page 74; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-329882/36.
                                                                                                                                                                                                                                                                                                                                                                            Bodnar JS,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABK71108
                                    02-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                              (REGC ) UNIV CALIFORNIA.
Antisense; interferon gamma
                    Human IFNGR2
                                                     AAD52338;
                                                                     AAD52338
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                                                                      standard;
                                                                                                                                  CCAAAGTGCTGGGATTA 403
                                                                                                                 CCAAAGTGCTGGGATTA
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                                                                                                                                                    Conservative
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                   antisense oligonucleotide,
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                                    (first entry)
                                                                      DNA;
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                                                                                                                                                            1.7%;
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, Wu C;
 receptor 2;
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Pred. No.
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  autoimmune disorder; cancer;
                    ISIS #142816.
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                                                                                                                                                                                                                                                                                                                         combined hyperlipidemia)
for diagnosing, treating
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ADA15247
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06-NOV-2003 ADA15247;

(first entry)

ADA15247

standard; DNA;

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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ##;
autoimmune thyroiditis; autoimmune insulinitis; multiple sclerosis;
diabetes; autoimmune arthritis; Crohn's disease; apoptosis; IFNGR2;
gene therapy; prophylaxis; human; phosphorothioate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotides for modulating Interferon gamma receptor 2, particularly useful for treating autoimmune disorders (e.g. multiple sclerosis or Crohn's disease), cancers or diseases caused by aberrant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI;
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                                                                                                                                                                          Seguence
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                                                                                                         Similarity
                                           CTCTGTCACCCAGGCTG
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CTCTGTCACCCAGGCTG
                                                                                                                                                                          BP; 3 A;
                                                                                       Conservative
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/mod_base= OTHER
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/note= "Phosphorothioate backbone; All
are 5-methylcytidines"
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                                                                                100.0%; --
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e= "2'-methoxyethyl
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                                                                                                                                                                       7 C; 5 G; 5 T;
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Pred. No.
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thes 0;
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                                                                                                                                DB 1; Length 20,
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CCDNA sequence, mouse HYPLIP1 genomic DNA, or the homologous human control the sequence is associated with a lipid disorder. Also claimed is an isolated polypeptide comprising a variant form of the mouse HYPLIP1 amino caid sequence, where the variant is associated with the lipid disorder. Also claimed is an cid sequence, where the variant is associated with the lipid disorder, an isolated polynucleotide having at least 12 contiguous nucleotides of the isolated polynucleotides, where the 12 contiguous nucleotides span caids of the encode polypeptides, where the 4 contiguous amino caids of the encode polypeptides, where the 4 contiguous amino caids span the variation position, a kit for the detection of the FCHLI cous comprising, an isolated antibody, identifying susceptibility to a lipid disorder which comprises comparing the nucleotide sequence of the suspected FCHLI allele with a wild-type FCHLI nucleotide sequence, where the difference between the suspected allele and the wild-type sequence identifies an altered HYPLIPI or FCHLI nucleotide sequence and a composition. Also disclosed is a transgenic animal which carries an altered hyPLIPI or FCHLI allele and a method for screening cand antibodies are useful for treation of FCHLI gene function as an anti-

card antibodies are useful for treation or reventing (e.g. gene therapy)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   allele; anti-lipid disorder; anti-cancer therapy; gene therapy; familial combined hyperlipidaemia; coronary artery disease; atherogenic lipoprotein phenotype; hyperapobetalipoproteinaemia; hyperrriglyceridaemia; low density lipoprotein subclass B; LDL; familial dyslipidemic hypertension; syndrome X; hypercholesterolaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel isolated polynucleotide comprising a mouse or human familial combined hyperlipidemia 1 gene having a variation that is associated a lipid disorder, useful for identifying susceptibility to the lipid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse HYPLIP1 locus
                                                                 and antibodies are useful for treating or preventing (e.g. gene therapy) a lipid disorder associated with expression of FCHIL, for diagnosis or prognosis of predisposition to lipid disorder, and cancer and for treating a lipid disorder such as familial combined hyperlipidaemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-SEP-2001; 2001US-00949428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US2003064372-A1
             coronary artery disease, atherogenic lipoprotein phenotype, hyperapobetalipoproteinaemia, hypertriglyceridaemia, low density lipoprotein (LDL) subclass B, familial dyslipidemic hypertension.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-540780/51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention discloses isolated polynucleotides comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ohmen
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TAFURI S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    insulin resistance; cancer; cytostatic; antilipaemic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ross D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          O
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human
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      The invention describes an isolated polypeptide (I) comprising a variant of form of a mouse HYPLIPI polypeptide sequence (SI) or a human FCHLI representation, where the variant form is associated with cancer, or an amino acid sequence having at least 65 % sequence identify to (SI) or (S2). A composition comprising DNA encoding (I) is useful for treating or preventing cancer associated with expression of FCHL1. FCHL1 gene or HYPLIP1 gene and its product are useful for the study of metabolic pathway and cellular mechanism to identify other genes, receptors and relationships that contribute to lipid disorder and cancer. FCHL1 gene or its fragments are useful in gene therapy to increase the amount of the expression products of the gene for the treatment of lipid disorder or cancerous cells. The sequence variation of FCHH1 gene or HYPLIP1 gene is also useful in the diagnosis and prognosis of predisposition to lipid disorder and cancer. Antisense
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse HYPLIP1
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                                                                                                                                                                                                             Claim 11; Page 37;
                                                                                                                                                                                                                                      Novel human FCHLI or mouse HYPLIPI polypeptide, screening, peptide therapy of lipid disorder or
                                                                                                                                                                                                                                                                                                        Ohmen
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polynucleotide sequences
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                                                                                                                         The invention comprises antisense oligonucleotides which are targeted to the human mdm2 gene. The antisense oligonucleotides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonucleotides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonucleotides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonucleotide of the invention. The present sequence contains 2'-methoxy-residues and has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense oligonucleotide; human; mdm2; hyperproliferation; hyperproliferative disorder; cancer; psoriasis; fibrosis; atherosclerosis; restenosis; apoptosis modulation; p21; ss; 2'-methoxy-residue; phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   expression used in the
                                                                                                                                                                                                                                                                               Novel antisense compound targeted to 5' untranslated region, coding region, or intron:exon junction of nucleic acid molecule encoding mdm2, useful for treating e.g. cancer, psoriasis or restenosis by inhibiting
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                                                                                                                                                                                                                                              Claim 4;
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Manoharan M;
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                                                                                                                The present sequence conta phosphorothicate backbone.
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              CTCTGTTACCCAGGCTG 951
 CTCTGTTACCCAGGCTG, 1
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AGTAGCTGGGACTACAG 745

Query Match Best Local S Matches 17

Similarity

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Mismatches

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1.5e+03; DB 1;

Indels

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Length 20;

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RESULT 1257
ABZ92717
The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory, steroid and ubiquinone. A composition of the invention CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an composition genstivity to adenosine, reducing levels of adenosine creceptor, producing bronchodilation, increasing levels of bully unione or CC lung inflammation, lung allergies, or a respiratory disease or condition. CC Note: The sequence data for this patent is not represented in the printed captification, but was obtained in electronic format directly from WIPO care from which was obtained in electronic format directly from WIPO care from which was obtained in electronic format directly from WIPO
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Miller
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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Sequence
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                                 ftp.wipo.int/pub/published_pct_sequences
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rion: lung allergy;
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RESULT 1258
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Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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lung inflammation; 1
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                                                                  l Similarity
17; Conserv
                      GGCTGGTCTCGAACTCC 225
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Tang L,
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                                                                                                                                                                        .int/pub/published_pct_sequences
                                                                    Conservative
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L, Shahabuddin
                                                                                                                                        A; 6
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                                                                                 1.7%;
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                                                                                                                                        G; 5 T; 0 U; 0 Other;
                                                                                    Score 17;
Pred. No.
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1 S;
                                                                      Mismatches
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                                                                                    1.5e+03;
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17

ATCTCGGCTCACTGCAA 983

<u>,</u>

Mismatches .

Indels

0,

Gaps

0

1.5e+03;

Local Similarity

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RESULT 1259
ABZ89862/c
ID ABZ8986
                                                                                 The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the cinitiation codon, coding region, 5' or 3' end genomic flanking regions, CC initiation codon, coding agion, 5' or 3' end genomic flanking regions, CC initiation codon, coding a polypeptide associated with lung and/or coding a polypeptide associated with lung and/or compassition of the invention and a second active agent comprising an cc antiinflammatory steroid and ubiquinone. A composition of the invention cc has antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also cf or enhancing the prophylactic or therapeutic respiratory effect of an cantiinflammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, increasing levels of ubiquinone or condition.
Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; constratic; gene therapy; antiaethmatic; through antiaethmatic; ensemble therapy; lung; adenosine sensitivity; antiense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABZ89862;
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                                                                                                                                         lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 5104; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-229219/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-APR-2001; 2001US-0286137P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; antisense; lung dysfunction; nasal airway dysfunction antiinflammatory steroid; ubiquinone; antiinflammatory; anti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human oligonucleotide sequence
                                                                             Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ubiquinone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
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                                                                                                                     ftp.wipo.int/pub/published_pct_sequences
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, Tang L,
                                                                               BP; 5 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
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L, Shahabuddin
                   1.7%;
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                     Score 17;
Pred. No.
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                                        DB 1;
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                                          Length
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ABD32
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CC oligonucleotides are derived from a gene encoding or regulating mRNA.

CC expression of a target polypeptide associated with lung airway or lung conversion of a cancer and can be anti-sense to the corresponding mRNA.

CC The invention also describes a kit, that comprises: (a) a delivery constructions for adding a carrier and for use of the kit. The composition of instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analysis of the addington composition of the kit. The composition of treating a respiratory, lung or malignant disease. The administered composition respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The compliance the administered conformation, allergies and/or bronchoconstriction and/or lung confilammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vascoonstriction, confilammation, emphysema, chronic obstructive pulmonary disease, pulmonary contents of the reduced adenosine content of the anti-sense oligos corresponding to the reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system of the oligonucleotides into products that free adenosine into the system.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             emphysema; chronic obstructive pulmonary disease; pulmonary transplantation rejection; ss; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-APR-2001; 2001US-0286036P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      surfactant depletion or hyposecretion, when administered to a mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes a novel composition (a) a first active agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 15; SEQ ID NO 14314;
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Tang
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L, Shahabuddin
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SSS & G
      RESULT 1261
ABD26092/c
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CC composition of igonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating composition, respiratory tract inflammation, allergies and composition adenosine sensitivity, levels of adenosine (A) or (A) receptors, composition or hyposecretion, when administered to a mammal. The collipse are derived from a gene encoding or regulating composition of a target polypeptide associated with lung airway or lung composition also describes a kit, that comprises: (a) a delivery composition also describes a kit, that comprises: (a) a delivery composition is eparate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, compassing a respiratory, immunosuppressive and cytostatic activity, is a beta adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition composition composition and is administered to reduce the production
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Best Local
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prevent
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Miller S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibros; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstrict respiratory distress syndrome; allergic rhinitis; pulmonary hypertens emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                  WPI;
                                                                                                                                                                                                                                       This invention describes a novel composition (a) a first active agent
                                                                                                                                                                                                                                                                                                                    Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-APR-2001; 2001US-0286036P
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                                                                                                                                                                                                                                                                      Claim 15;
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17; Conserv
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Tang
                                                                                                                                                                                                                                                                      SEQ ID NO
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Shahabuddin
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                                                                                                                                                                                                                                                                      5104; 763pp;
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Pred. No
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        fibrosis;
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RESULT 1262
ABD28947
ID ABD2894
XX ABD2894
XX ABD2894
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XX ABD2894
XX Person New Yespira
KW respira
KW pulmona
XX Homo sa
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XX Claim
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20
       This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) recepto
                                                                                                                                                                                                                                                                            Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N58473-derived oligonucleotide SEQ ID 7959.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JUL-2004
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                                                                                                                                                                                                                                                   bronchodilating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EPIGENESIS PHARM INC
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Pred. No.
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1.5e+03;
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       allergies and (A) or (A) receptors,
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RESULT 1263
ADJ60957
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XX ADJ6095
XX Oligonu
XX interle
KW interle
KW airway
KW eystic
KW gulmone
KW gulmone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC oligonuclectides are derived from a gene encoding or regulating CC expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition CC of the invention has antiallergic, antiinflammatory, antiasthmatic, CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered CC composition comprises oligo and is administered to reduce the production cor availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The CC inflammation, allergies and/or bronchoconstriction and/or lung conflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, conflammation, emphysema, chronic obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary cancer.
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Best Local Sim:
Matches 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
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                                                                                         WPI; 2004-203534/19.
                                                                                                                                                  Shahabuddin
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it any unwanted
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in S,
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                                                                                                                                               Sandrasagra
H, Cong H;
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Pred. No.
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Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes

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e.g.,

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RESULT 1264
ADL32184
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to an oligonucleotide anti-sense to e.g., cc initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-cc end of nucleic acid target comprising gene(s) chosen from e.g. cc interleukin (IJ)-4 receptor, IL-5 receptor or salts of the coligonucleotide and optionally surfactant operatively linked to the coligonucleotide. The method is useful for preventing or treating a cc oligonucleotide. The method is useful for preventing or treating a cc of a subject an effective amount of an inhibitor. The oligonucleotide is cc useful for production of a medicament for the prevention and/or treatment cc of a respiratory or lung disease. The respiratory or lung disease is cc chosen from airway inflammation, allergy(ies), asthma, impeded cc respiration, cystic fibrosis (EC), chronic obstructive pulmonary diseases (CODD), allergic rhinitis (AR), acute respiratory distress syndrome cc (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway construction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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11-JAN-2000;
02-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                    human; medicine; signal transduction; glycoprotein; oligo-capping method; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                              Clone specific PCR primer to
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This invention relates to a novel primers useful for synthesising full length cDNA molecules that encode human proteins. Specifically, it refers
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                                                                New oligonucleotide primers length human cDNAs.
                                                                                                                                             Ota T,
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les 17; Conserv
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se e.g.,
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                                        18;
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2000JP-00183865.
2000EP-00114089.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                               Isogai T,
a T, Nagai
                                      4217; 1340pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 17; pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                              (830 cDNAs) useful
                                                                                                                                                                                                                                                                                                                                                                                                              amplify human full length cDNA SeqID 4217.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 T; 0 U; 0 Other;
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                                                                                                                                Hayashi K,
K, Kojima
                                      English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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                                                                                                                                Ishii S,
S, Otsuki
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                                                                              synthesizing
                                                                                                                                                                                                                                                                                                                                                                                      transcription,
                                                                                                                               Kawai
T, Ko
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length

isoprenylcysteine carboxyl methyltransferase

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RESULT 1265
ADJ10599
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    to secretory or membrane proteins that are potential therapeutic agents/
target molecules in the field of medicine, and in particular genes
encoding proteins that are associated with signal transduction,
glycoproteins and transcription. The present invention describes a method
for efficiently cloning a full length human cDNA from both the 5' and 3'
ends using the oligo-capping method. This oligonucleotide sequence is a
human clone specific PCR primer used in an exemplification of the
         This invention relates to a novel antisense compounds that modulate the expression of isoprenylcysteine carboxyl methyltransferase (also known as ICMT, PCCMT, pcMTase, PPMT, PPMTase, HSTE14, MST098 and MST098) and located on chromosome 1p36. Specifically, it refers to compositions useful for inhibiting the expression of isoprenylcysteine carboxyl methyltransferase, which normally participates in cellular events such as growth factor signal transduction, cell replication, vesicular transport and the post-translational modification of the Ras family of GTPases. The present invention describes antisense oligonucleotides that comprise at least one modified sugar moiety, a 2'-O-methoxyethyl (2' MOE) and at least one modified mucleobase, a 5-methylcytosine. Accordingly, these compounds are useful for treating a disease or condition associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PPMT; PPMTase; HSTE14; MST098; MSTP098;
growth factor signal transduction; cell replication; vesicular transport;
hyperproliferative disorder; cancer; inflammatory; hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20
                                                                                                                                                                                                                                                              nucleic acid encoding isoprenylcysteine carboxyl methyltransferase, useful for treating cancer, hypertension, or cardiovascular or
                                                                                                                                                                                                                                                                                                                                                                                                                             31-MAY-2002; 2002US-00159834
                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-MAY-2002; 2002US-00159834.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cardiovascular; cytostatic; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADJ10599;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADJ10599 standard; DNA;
                                                                                                                                                                                                                   Example 15; SEQ ID NO 126; 62pp; English.
                                                                                                                                                                                                                                                    inflammatory
                                                                                                                                                                                                                                                                                                 New compounds,
                                                                                                                                                                                                                                                                                                                                  WPI; 2004-081071/08
                                                                                                                                                                                                                                                                                                                                                                 Dobie
                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   isoprenylcysteine carboxyl methyltransferase;
PPMTase; HSTE14; MST098; MSTP098;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA oligo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                    disease.
                                                                                                                                                                                                                                                                                                                                                                                                PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                 particularly antisense oligonucleotides targeted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ss; PCCMT; pcMTase;
                                                                                                                                                                        (also known as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>.</u>.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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RESULT 1266
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ស្ត្រិនិង្គិនិង្គិនិង្គិនិងិ
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antiinflammatory, hypotensive and cardiant activities and are useful for research reagents and in diagnostics. This oligonucleotide sequence is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disorder (e.g. cancer), an inflammatory condition, hypertension cardiovascular disease. As such, they exhibit cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA oligo representing a preferred target site for antisense the human isoprenylcysteine carboxyl methyltransferase, given in an
                            New compounds, particularly antisense oligonucleotides targeted to nucleic acid encoding isoprenylcysteine carboxyl methyltransferase, useful for treating cancer, hypertension, or cardiovascular or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hyperproliferative disorder; cancer; inflammatory; hypertension; cardiovascular; cytostatic; antiinflammatory; hypotensive; cardiant; ICMT; antisense; phosphorothioate backbone; 2' MOE wing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PPMT; PPMTase; HSTE14; MST098; MSTP098;
growth factor signal transduction; cell replication; vesicular transport;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADJ10546 standard; DNA; 20
                                                                                                                                                                                                                                                                                                           31-MAY-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Phosphorothioate antisense DNA oligo to modulate human ICMT SeqID 73.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US2003228688-A1
                                                                                                                                           2004-081071/08
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17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    isoprenylcysteine carboxyl methyltransferase; ss;
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                                                                                                                                                                                                                                                                                                           2002US-00159834.
                                                                                                                                                                                                                                                        PHARM INC
                                                                                                                                                                                                                                                                                                                                                                  2002US-00159834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides.
cytidine nucleobases are 5-methylcytidine."
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
/note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides.
cytidine nucleobases are 5-methylcytidine."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag=
/mod_ba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        *tag=
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>,</u>
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This invention relates to a novel antisense compounds that modulate the expression of isoprenylcysteine carboxyl methyltransferase (also known as ICMT, pCCMT, pCMTase, PPMT, pPMTase, HSTE14, MST098 and MSTP098) and CC located on chromosome 1936. Specifically, it refers to compositions cuseful for inhibiting the expression of isoprenylcysteine carboxyl cystein factor signal transduction, cell replication, vesicular transport cystem factor signal transduction, cell replication, vesicular transport cystes and the post-translational modification of the Ras family of GTPases. The present invention describes antisense oligonucleotides that comprise at CC least one modified sugar moiety, a 2-O-methoxyethyl (2', MOE) and at CC compounds are useful for treating a disease or condition associated with isoprenylcysteine carboxyl methyltransferase such as a hyperproliferative CC disorder (e.g. cancer), an inflammatory condition, hypertension or CC cardiovascular disease. As such, they exhibit cytostatic, antiinflammatory, hypotensive and cardiant activities and are useful for research reagents and in diagnostics. This oligonucleotide sequence is a phosphorothioate antisense DNA oligo used to modulate human carboxyl methyltransferase expression in an
                                                                                                                                    Matches
                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                              Sequence 20 BP; 5 A; 5 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 15; SEQ ID NO 73; 62pp; English
                                                                                                                                                                                                                                                                                                                                    exemplification of the invention
                                                              385
17
                                                                                                                                    17;
                                                                                                                                                                Similarity
                                      TCCCAAAGTGCTGGGAT 401
TCCCAAAGTGCTGGGAT 1
                                                                                                                                 Conservative
                                                                                                                                                                100.0%;
                                                                                                                                    0
                                                                                                                                                                   Score 17;
Pred. No.
                                                                                                                                    Mismatches
                                                                                                                                                                                                   DB 1;
                                                                                                                                                                   1.5e+03
                                                                                                                                    0
                                                                                                                                                                                               Length 20;
                                                                                                                                    Indels
                                                                                                                                    <u>..</u>
                                                                                                                                    Gaps
                                                                                                                                    0
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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
Homo sapiens.
Synthetic.
                                                                                                             Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADM15229 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1416.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADM15229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20
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modified\_base

residues

mod.

base= OTHER

note= "phosphorothioate linkages residues are 5-methylcytidines"

and

modified\_base

Location/Qualifiers

modified\_base

/note= " 16. .20

mod

\_base= e= "2'

OTHER

-O-methocyethyls"

\*tag=

/\*tag= /mod\_ba

\_base= e= "2'

OTHER

-O-methoxyethyls"

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AD046446
AD04
XX
AC AD04
XX
XX
DT 15--:
XX
Huma
XX
Huma
XX
Huma
XX
Hum
XW
LTY,
XW
Air
XW
Air
XW
Air
XX
AN
OS
Homm
XX
Home
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to CC 9334.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC maying a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, antiarthritic, vasotropic, CC ophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or CC condition associated with mpGES-1 a disease or CC condition associated with mpGES-1 and can be used for preparing a composition for treating a disease or CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 1268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2004028458-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compound, having a sequence targeted to a nucleic avencoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gierse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-SEP-2003; 2003WO-US030374.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-APR-2004
                                                                         Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD33; ICAM; VCAM; tryptase a; tryptase b; PD54 A; PD54 B; PD54 C; PD54 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation, inflammatory disease; airham; lung allergy; inflammation, inflammatory disease; corporation; cystic fibrosis; CF chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                            ADO46446 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; SEQ ID NO 1416; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ischemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-305094/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-SEP-2002; 2002US-0413549P
                                                                                                                                                                                                                                                                 Human oligonucleotide #1812.
                                                                                                                                                                                                                                                                                                            15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          769
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TTTTTGTATTTTTAGTA 785
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 13 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.7%;
llarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                            (first entry)
                                                                bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 C; 0 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     μ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 17; DB Pred. No. 1.5
                                                              airway obstruction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1; Lo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                bronchoconstriction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    a nucleic acid
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ADP08719
ID ADP0
XX
AC ADP0
XX
DT 26-A

ADP08719;

ADP08719 standard; DNA;

20

0

26-AUG-2004

(first entry)

RESULT 1269

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                           S
                                                                                                                                                                                CC codon, coding region, 5' or 3' intron-exon junction, intron or region cod with 2-10 nucleotides of the 5' end or 3' end of a nucleic acid target chosen from a gene encoding interleukin (II) 4 receptor, interleukin (II) 6-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention compounds that binds to one or more nucleic acid target(s) or expressed product(s), for the compound encoding interleukin or expressed product(s), for the coligonucleotides are useful for reducing or inhibiting expression of a coligonucleotides are useful for reducing or inhibiting expression of a coligonucleotides are useful for reducing or inhibiting expression of a coligonucleotides are useful for reducing or inhibiting expression of a compound that binds to coligonucleotides are useful for preventing or treating or inhibiting expression of a compound that binds to coligonucleotides are useful for preventing or treating or inhibiting expression of a conspiratory or lung disease is associated with hyper-responsiveness to compound that binds are compound that binds are compound that binds are conspiratory or lung disease to inflammation or an inflammatory disease. The respiratory or lung disease to compound that and/or inflammation, allergy, asthma, impeded respiration, constitution or an inflammation, bronchitis, airway obstruction or invention or compound inflammation, bronchitis, airway obstruction or invention. This sequence represents an oligonucleotide of the invention.
                                                                Matches
                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nyce JW, Sandrasagra A, Shahabuddin S, Lu H, C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-APR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-JUL-2003; 2003US-00627930
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2004049022-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to oligonuclectides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 2; SEQ ID NO 1813; 174pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-293804/27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SAND/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-APR-2002;
                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (/HHn7
                                 209
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SANDRASAGRA A.
TANG L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SHAHABUDDIN S
                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CONG H.
                               GGCTGGTCTCGAACTCC 225
                                                                                                                                      20
GGCTGGTCTCGAACTCC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002WO-US013135
2002WO-US013143
                                                                   Conservative
                                                                                                                                      BP; 4
                                                                                                                                      Ą
                                                                                  100.0%;
                                                                                                                                      6 C; 5
                                                                  1.7%; Score 17; DB
100.0%; Pred. No. 1.
ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tang
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                                                                                                                                      G;
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                                                                                                                                      T; 0
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                                                                                                                                       U; 0 Other,
                                                                     1.5e+03;
                                                                                                     DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Miller
                                                                     0;
                                                                     Gaps
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RESULT 1270
ADP0928
ID ADP0928
XX
AC ADP0928
XX
DT 26-AUG-
XX
DT 26-AUG-
XX
DT Extend
XX
Extend
XX
Secretc
XW breast
XW secretc
XW single
XX
Homo sa
XX
PN WO2004(
XX
PF 25-NOV-
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                                                                                                                                                                                                                                                                                                                            뭐
                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel method for identifying a subject at risk of breast cancer which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a risk of breast cancer, as well as therapeutic and prophylactic treatments that specifically target breast cancer, such as gene therapy. The current sequence is that of an Extend primer of the invention which was used to genotype single nucleotide polymorphisms within human glycoprotein VI (platelet) (GP6; GPIV;GPVI) DNA which is located at chromosomal position 19q13.4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying a subject at risk of breast cancer by detecting or absence of one or more nucleotide polymorphic variations, diagnosing, preventing and/or treating breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Roth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              single nucleotide polymorphism.
                                                                                                           breast cancer; cytostatic; gene therapy; human; chromogranin B; secretogranin 1; SCG1; chromosome 20pter-p12; ss; PCR; primer; single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-NOV-2002; 2002US-0429136P
24-JUL-2003; 2003US-0490234P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2004047767-A2
 25-NOV-2003; 2003WO-US037966
                             10-JUN-2004
                                                       WO2004047767-A2
                                                                                                                                                                                                                           ADP09281
                                                                                                                                                                                                                                                      ADP09281 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-NOV-2003; 2003WO-US037966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-JUN-2004
                                                                                                                                                                                              26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SEQU-) SEQUENOM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gt cancer;
GPIV; GPV
                                                                                                                                                                                                                                                                                                                                                      967
                                                                                                                                                                                                                                                                                                                                                                                17; Conserv
                                                                                                                                                                   primer
                                                                                                                                                                                                                                                                                                                            4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3; Page 83; 286pp; English.
                                                                                                                                                                                                                                                                                                                            ATCTCGGCTCACTGCAA 20
                                                                                                                                                                                                                                                                                                                                                    ATCTCGGCTCACTGCAA 983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nelson MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GPVI;
                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 5
                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                              (first entry)
                                                                                                                                                                   76 used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                      A,
                                                                                                                                                                                                                                                                                                                                                                                            1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                      6 C; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ូ
                                                                                                                                                                   to genotype human chromogranin B polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Braun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genotype human glycoprotein VI polymorphism
                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene therapy; human; platelet glycoprotein VI; e.19q13.4; ss; PCR; primer; SNP;
                                                                                                                                                                                                                                                                                                                                                                                  0:
                                                                                                                                                                                                                                                                                                                                                                                                                                      G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Þ
                                                                                                                                                                                                                                                                                                                                                                               Score 17; DB Pred. No. 1.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kammerer
                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Le
1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Reneland
                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the presence, useful for
                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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RESULT 1271
AAX86419
ID AAX8641
XX
AC AAX8641
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel method for identifying a subject at risk of breast cancer which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a risk of breast cancer, as well as therapeutic and prophylactic treatments that specifically target breast cancer, such as gene therapy. The current sequence is that of an Extend primer of the invention which was used to genotype single nucleotide polymorphisms within human chromograpin B (CHG); secretogranin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diagnosing, preventing and/or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-NOV-2002; 2002US-0429136P.
24-JUL-2003; 2003US-0490234P.
                                                                                                                                                                                                                                                               Homo
                                                                                                                                                                                                                                                                                                                                                                     29-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 5; Page 103; 286pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Identifying a subject at risk of breast cancer by detecting the presence or absence of one or more nucleotide polymorphic variations, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-441082/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Roth RB, Nelson MR,
                                                  MMSC1, an MMAC1 (tumor suppressor) interacting protein and
                                                                                                    Bartel PL,
                                                                                                                                                        20-JAN-1998;
                                                                                                                                                                                  19-JAN-1999;
                                                                                                                                                                                                           22-JUL-1999
                                                                                                                                                                                                                                     WO9936566-A1
                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                      MMAC1
                                                                                                                                                                                                                                                                                                                 Human; MMSC1
                                                                                                                                                                                                                                                                                                                                           PCR primer PDZK5.6P used to amplify DNA encoding MMSC1 protein.
                                                                                                                                                                                                                                                                                                                                                                                               AAX86419;
                                                                                                                                                                                                                                                                                                                                                                                                                       AAX86419 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1;SCG1) DNA which
                                                                                                                             (MYRI-) MYRIAD GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SEQU-) SEQUENOM INC
                                                                                                                                                                                                                                                             sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                390
                                                                             1999-458472/38
                                                                                                                                                                                                                                                                                                  ; MMSC1 protein; MMAC1 interacting protein; tumour suppression; pathway; immunogen; cancer; cell neonlastic tumour suppression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ,_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAGTGCTGGGATTACAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 BP; 7 A; 2 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAGTGCTGGGATTACAG
                                                                                                    Tavtigian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.7%;
ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                                        98US-0071861P
                                                                                                                                                                                  99WO-US000995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      is located at chromosomal position 20pter-p12.
                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                      ٧S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Braun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 17;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kammerer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; L
. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Reneland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₽,
                                                                                                                                                                                                                                                                                                     PCR primer; ss.
                                                     related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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Example 5; Page 51; 107pp; English

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CC MMSC1 protein. The PCR templates were derived from tumour cell lines, and CC MMSC1 protein. The PCR templates were derived from tumour cell lines, and CC the amplicons were tested for mutations. The MMSC1 protein is a MMAC1 CC interacting protein which is involved in tumour suppression activity in CC the MMAC1 pathway. MMSC1, antigenic fragments or fusion proteins of these CC are used as immunogens for antibody production. Primers derived from CC MMSC1 genomic clones can be used for identification of MMSC1 genes and CC for synthesis by amplification of MMSC1 DNA or RNA. Detecting an CC alteration in MMSC1 can be used to diagnose cancer. A germline alteration CC in an MMSC1 gene is indicative of a predisposition to cancer. A somatic CC contaction in an MMSC1 gene is indicative that the tissue is cancerous. CC interactions can be used for detection of alterations in MMAC1 associated CC with cancer. Wild-type MMSC1 or a homologue can be used to supply wild-cype MMSC1 gene function (or a substantially similar function) to a cell, which has lost the gene function due to a MMSC1 gene mutation. The gene CC suppresses neoplastic growth of the cell. Transgenic animals having an CC useful in treating cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP specific upper PCR primer SEQ ID 1929.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAH39133 standard;
                                                                                                                                                                                                                                                                                                                                                                 15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                              13-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                         New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                  WPI; 2001-290930/30
Claim 1; Page 59; 83pp; English.
                                                                                                                                                                                                                                        Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                        (ORCH-) ORCHID BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             635 CTCTGTCACCCAGGCTG
                                                            sampie.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTCTGTCACCCAGGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 2 A; 8 C; 5 G; 6 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                              2000WO-US028436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                    99US-0160096P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                        Pohl M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 17;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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includes kits for determining the presence or absence of a SNP, using the CC oligonucleotides of the invention. The PCR primers are used to amplify a CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or CC identity of a SNP and for genotyping nucleic acid samples for e.g. to cases by association analysis the genotype of an individual or group of CC individuals, having a pathological phenotypic trait suspected of being CC caused by one or more SNPs. Phenotypic traits include diseases e.g. CC agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular CC dystrophy, familial hypercholesterolaemia, polycystic kidney disease, cC craits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune CC disease of which a component is or may be genetic such as autoimmune concerning including, rheumatoid arthritis, multiple sclerosis, concerning the method is also useful in forensic investigations and conternity analysis. The present sequence represents a PCR primer specific for a human SNP containing DNA sequence
Sequence 21 BP; 4 A; 7 C; 5 G; 5 T; 0 U; 0
                                                                                                                            for a human SNP containing DNA sequence
                  Other;
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88888888888888888888888888888888

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RESULT 1273
AAF23445/c
                                                                                        Matches
                                                                                                    Query Match
Best Local
                                                            207 CAGGCTGGTCTCGAACT 223
                                                                                        17;
                                              N
                                                                                                  Similarity
                                              CAGGCTGGTCTCGAACT 18
                                                                                          Conservative
                                                                                                    100.0%;
                                                                                        <u>.</u>
                                                                                                     Score 17; DB 1; Length 21; Pred. No. 1.6e+03;
                                                                                           Mismatches
                                                                                          1.6e+03;
                                                                                          Indels
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immune disorder; neurodegenerative disease; allergic reaction; respiratory problem; organ transplantation; contraceptive; human; PCR primer; proliferative disorder; ss.
                                                           SECX; secreted protein; cancer; angiogenesis; wound healing; immune disorder; neurodegenerative disease; allergic reactio
                                                                                                                                               Forward PCR primer for amplification of DNA encoding
                                                                                                                                                                                                                                                                AAF23445;
                                                                                                                                                                                                                                                                                                                      AAF23445 standard; DNA;
                                                                                                                                                                                                           20-MAR-2001
                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                         21
                                                                                                                                                                                                                                                                                                                            ВЪ
                                                                                                                                                     SEC3
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Synthetic

WO200070046-A2

23-NOV-2000

12-MAY-2000; 2000WO-US013291.

14-MAY-1999; 12-JAN-2000; 10-MAR-2000; 2000US-0175744P. 2000US-0188274P. 2000US-00569269. 99US-0134315P

11-MAY-2000;

(CURA-)

CURAGEN CORP

₽, Fernandes Έ Boldog

WPI; 2001-025020/03.

New SECX polypeptides and nucleic acids useful for treating or preventing cancer, other disorders related to angiogenesis, neurodegenerative diseases, autoimmune disorders and allergic reactions.

Example 8; Page 117; 132pp; English

Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention

Polynucleotide sequences AAF23410 - AAF23419 encode secreted SECX

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AAI78386/c
ID AAI783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1274
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            proteins AAB49649 - AAB49658. Sequences AAF23420 - AAF23450 represent primers and probes used in the isolation and characterisation of the SECX DNA sequences of the invention. The new polypeptides and nucleic acids can be used in screening assays, detection assays, preventive or predictive medicine, therapeutic and prophylactic treatment, and pharmacogenomics. Specifically, the SECX polypeptides and nucleic acids are useful for treating cancer; other disorders related to angiogenesis e.g. abnormal wound healing, psoriasis; neurodegenerative diseases; immune disorders; liver cirrhosis; benign tumours; fibrocystic conditions and tissue hypertrophy (e.g. benign prostatic hypertrophy); allergic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        parasites, and as a vaccine. SECX antibodies may be used to isolate or detect SECX proteins, monitor protein level in tissue as part of a clinical testing procedure, treat proliferative disorders including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 reactions and conditions such as asthma and other respiratory problems; as well as in treating or preventing diseases associated with organ transplantation, atherosclerosis-associated diseases or disorders. The polypeptides can also be used for bone, cartilage, tendon, ligament and/or tissue growth or regeneration, wound healing, tissue repair and replacement, gut protection or regeneration, as a contraceptive, to inhibit thromboses, infections caused by bacteria, virus, fungi and other inhibit thromboses, infections caused by bacteria, virus, fungi and other inhibit thromboses.
   AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53329 represent peptides related to human polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 4 A; 7 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-NOV-1999;
29-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human silent SNP containing nucleic acid SEQ:5327.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAI78386 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tumours and
                                                                                                                                       Claim
                                                                                                                                                                                                                                 Polymorphic nucleic acid sequences,
                                                                                                                                                                                                                                                                                               WPI; 2001-356160/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-NOV-2000; 2000WO-US032758
                                                                                                                                                                                                                                                                                                                                                                                                                             (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     731 TAGCTGGGACTACAGGC 747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21
                                                                                                                                    Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TAGCTGGGACTACAGGC 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             benign hyperplasias
AAM53329 represent peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0168138P.
2000US-00726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                    2140;
                                                                                                                                                                                                                                                                                                                                                              Leach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.7%;
                                                                                                                                                                                                                                                                                                                                                              3
                                                                                                                                    2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     В₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 17;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                 useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1; L:
1.6e+03;
                                                                                                                                                                                                                                 ä
                                                                                                                                                                                                                             genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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can be used in

gene

protein

Query Match Best Local

Sequence 20

BP; 5 A; 5 C; 6 G; 4 T; 0 U; 0 Other;

correct

P

field.)

Matches

18;

Conservative

0

Similarity

1.7%;

Score 16.8; D Pred. No. 1.6e 0; Mismatches

.6e+03

16.8;

DB 1;

Length

20;

Indels

<u>.</u>

Gaps

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC thermapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC example, (I) may be used to treat disorders by rectifying mutations or CC deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC therefore which patients may be in need of restorative therapy. The CC polypeptides encoded by (I) may be used as antigens in the production of CC antibodies specific for polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic CC polypeptides in samples
RESULT 1275
AAQ53171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                         The sequence is that of a primer specific for the D9S58 marker polymorphism which may be used in the detection of a gene associated with familial dysautonomia (FD). It may be used in a test kit for screening of foetuses and individuals at risk through their family. (Updated on 25-MAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
09-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 51 BP; 9 A;
                                                                                                                                                                                                                              Blumenfeld A,
                                                                                                                                                                                                                                                                                           29-MAY-1992;
16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                       09-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Probe; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Familial dysautonomia detection D9S58 primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ53171 standard;
                                                                                                                     Disclosure; Page
                                                                                                                                                 Detection of a gene associated with familial dysautonomia -
human chromosome 9 for DNA polymorphism linked to the gene.
                                                                                                                                                                                                  WPI; 1993-405845/50.
                                                                                                                                                                                                                                                                                                                                        25-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                                                                  WO9324657-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ53171;
                                                                                                                                                                                                                                                             (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                472 AGGATGAAGTGCAGTGGTGATCACAGCTCACTGCAGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGGTTGCAGTGACCCGGGATCGTGCCACTTCACTCCAGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chromosome
                                                                                                                                                                                                                                Gusella JF,
                                                                                                                                                                                                                                                                                          92US-00890719
93US-00049678
                                                                                                                                                                                                                                                                                                                                        93WO-US004946
                                                                                                                     25; 50pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17 C; 16 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9; FD;
                                                                                                                      English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                Breakefield
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 17;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene; screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
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RESULT 1276

RAQ47775

ID AAQ47775

AC AAQ4777

AC Sixteer

CC CAAQ4777

CC AAQ4777

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                                                                                                                                   RESULT 1277
AAQ63001/c
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SARAKE
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                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense oligo-nucleotide(s) to nucleolar protein genes - used diagnosis and treatment of hyperproliferative disease, partic. malignancies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BAYU )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cell proliferation-associated protein; p120; nucleolar protein; malignant cell growth; inhibition; hyperproliferation; disease; human malignancy; breast cancer; 120 kDa nucleolar protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ47775;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ47775 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sixteen oligonucleotide sequences (AAQ47764-Q47779) were designed based on different regions of the sequence coding for the nucleolar protein p120, associated with hyperproliferative diseases. Those antisense oligonucleotides directed to the 3'-untranslated region were found to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human mailgnamey; brown co
3'-untranslated region; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
23-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                      have particular inhibitory activity. Oligonucleotides AAQ47772 and AAQ47777 have demonstrated high activity in inhibiting a number of cancers. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-JAN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-SEP-1993
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                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 13; Page 32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-FEB-1992;
                    25-MAR-2003
17-NOV-1994
                                                                             AAQ63001;
                                                                                                                     AAQ63001 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          725
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                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCTGAGTAGCTGGGACTACA
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                                                                                                                                                                                                                                                           TCTCGAACTCCCGACCTCAG 234
                                                                                                                                                                                                                     TCTCGAACACCTGACCTCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
(first en
                                                                                                                                                                                                                                                                                                     Conservative
                  (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   50pp; English.
                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                   1.7%;
                                                                                                                                                                                                                                                                                                                                                                                8 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Perlaky L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
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                                                                                                                       BP
                                                                                                                                                                                                                                                                                                   Score 16.8; D
Pred. No. 1.6e
O; Mismatches
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                                                                                                                                                                                                                                                                                                                        .6e+03
                                                                                                                                                                                                                                                                                                                                        DB 1; Length 20;
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AAQ75579
                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                            The sequences given in AAQ63001-02 are primers which were used to compare linkage between a predisposition to hypertension with the angiotensin-converting enzyme (ACE) gene. These primers map to the 5'region or the exons of the AGT gene. The amplified products are analysed by single stranded conformation polymorphisms (SSCP) to identify any differences which were then sequenced and compared to the normal gene. These primers can especially be used to determine a predisposition to essential hypertension or pregnancy-induced hypertension. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer; polymerase chain reaction; PCR; amplify; angiotens:
predisposition; hypertension; human; 5' region; exon;
single stranded conformation polymorphism; SSCP; essential
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pregnancy-induced hypertension; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Use of angiotensinogen gene variants - for determine to hypertension, partic essential hypertension or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-APR-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hypertension/ACE linkage analysis primer
                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Page 23; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9408048-A1
                                                                                   Synthetic
                                                                                                                                              Reverse transcription primer used in cDNA analysis technique
                                                                                                                                                                         04-AUG-1995
                                                                                                                                                                                                AAQ75579;
                                                                                                                                                                                                                        AAQ75579 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1994-135608/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lalouel J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UTAH ) UNIV UTAH RES FOUND.
(INRM ) INSERM INST NAT SANTE & RECH MED.
             16-APR-1993;
                                     01-NOV-1994.
                                                             JP06303997-A
                                                                                                          aggregate;
                                                                                                                      Analysis; gene
                                                                                                                                                                                                                                                                                                              641 CACCCAGGCTGGAGTGCAGT 660
                                                                                                                                                                                                                                                                                      20
                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                        CTCCGAGGCTGGAGTGCAGT 1
                                                                                                             restriction enz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jeunemaitre X,
                                                                                                                                                                                                                                                                                                                                                                                       BP; 4 A;
                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     92US-00952442
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             93JP-00112515
                                                                                                                                                                                                                                                                                                                                                 1.7%;
                                                                                                                                                                                                                                                                                                                                                                                       8 C;
                                                                                                             enzyme;
                                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                       5 G; 3 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lifton RP,
                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                     Score 16.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                    Pred.
                                                                                                                                                                                                                                                                                                                                     ed. No. 1.6e+03
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                        0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for determining a predisposition rtension or pregnancy-induced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Soubrier F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amplify; angiotensinogen; AGT;
                                                                                                                                                                                                                                                                                                                                                                                         0 Other;
                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                          primer; cDNA;
                                                                                                                                                                                                                                                                                                                                         Indels
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AAQ75581
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 5; 11pp; Japanese.
                                        A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files Ap075547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Analysis of cDNA and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-APR-1993;
                                                                                                                                                                                                                                                                                                                                16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ75581 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NITE ) NIPPON TELEGRAPH & TELEPHONE
                                                                                                                                                              Disclosure; Page 5; 11pp; Japanese.
                                                                                                                                                                                            Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                   16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                              01-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reverse transcription
                                                                                                                                                                                                                                                                       (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              alysis of cDNA and gene expression - digestion with restriction enzymes.
                                                                                                                                                                                                                                         1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               429
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene expression; reverse
; restriction enzyme; ss.
                             Бe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 2 A; 0 C; 1 G; 17 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                93JP-00112515
                            to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               primer used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP
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Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  transcription; primer; cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
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Sequence 20

BP; 2 A;

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G; 18 T; 0 U; 0 Other;

RESULT 1281

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RESULT 1280
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ID AAT4114
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Best Local S
Matches 18
                                                                    Query Match
Best Local
                                                       Matches
                                                                                                                                      Primers T41001-T41382 are derived from novel human gene signature (GS) sequences which did not match with sequences deposited in Genbank release 76. The GS sequences (T19001-T26837) were obtained from 3'-directed cDNA libraries prepared from various human tissues; synthesis of cDNA was initiated from the 3'-end of mRNA by using poly(T) as the sole primer. Each library is constructed so as to reflect accurately the relative abundance of different mRNAs in the particular tissue from which it was derived. The appearance frequency of a given GS in a cDNA library can be determined (esp. using primers and probes derived from the GS sequences) as a means of diagnosing abnormal cell function or for recognising different cell types. The primers T41147-8 amplify clone pm0368 which comprises the GS HUMGS001308 (T20308), located on chromosome 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gene signature; messenger RNA; mRNA; relative abundance; frequency; human; cloning; mapping; non-biased library; diagnosis; detection; cell typing; abnormal cell function; primer; PCR; amplification;
                                                                                                                                                                                                                                                                                                                                                            Single-stranded DNA for identifying gene signatures - isolated from 3'-directed human cDNA library that reflects relative abundance of corresp. mRNA in specific human tissues.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matsubara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polymerase chain reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human gene signature HUMGS01308-derived sense primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT41147
                                                                                                              Sequence 20 BP; 6 A; 4 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                 Example 7;
                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1995-206931/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-NOV-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MATS/) MATSUBARA K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             595
                          387
                                                       18;
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18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OKUBO
                                                                      Similarity
                          CCAAAGTGCTGGGATTACAG 406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TTTTTTTTTTTTTTTAAT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TTTTATTTTATTTTAAT 614
CCAAAGTGCTAGGGTTACAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                 7,
                                                                                                                                                                                                                                                                                                                                 Fig 7; 2245pp;
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                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Okubo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93JP-00355504.
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                                                                    1.7%;
                                                                                                                                                                                                                                                                                                                                  Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88
                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                      Pred. No.
                                                                                   Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 16.8; DB 1;
Pred. No. 1.6e+03;
20
                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                   16.8;
                                                                       .6e+03
                                                                                   DB 1;
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                                                                                   Length
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AAT18321/c
ID AAT183
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AC AAT183
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XX
OS-JUN
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BRCA1;
XW BRCA1;
XW BRCA1;
XW Chromo
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Chromo
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Chromo
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II-AUC
PR 11-AUC
PF 11-AUC
PF 11-AUC
PR 29-AUA
PR 29-AUA
PR 07-JUN
RESULT 1282
AAT17537/c
ID AAT1753
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AC AAT1753
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DT 03-OCT-
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Best Local S
Matches 18
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02-SEP-1994;
16-SEP-1994;
29-NOV-1994;
29-NOV-1995;
07-JUN-1995;
07-JUN-1995;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BRCA1; breast cancer; ovary cancer; predisposing gene; susceptibility gene; diagnosis; prognosis; gene therapy; chromosome 17q; primer; polymerase chain reaction; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 Kindred families provided genetic evidence for localisation of the human cancer predisposing gene, BRCA1 (see AAT18310), to a sufficiently small region of 17q for the appln. of positional cloning strategies. 15 short tandem repeat markers assayable by PCR were used to refine this localisation. Primer sequences for 4 of these markers were AAT18315-16 for DS178754, AAT18317-18 for DS178975, AAT18319-20 for tdj1474, and AAT18312-22 for tdj1239. The region contg. BRCA1 was estimated to be approx. 650 kb and to be flanked by tdj1474 and U5R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Skolnick MH, Goldge
Shattuck-Eidens DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JUN-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated human cancer predisposing gene, BRCA1 - used to develop prods. for diagnosis, prognosis and therapy of cancers, partic. breas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1996-139703/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-AUG-1995;
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                            03-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                         AAT17537;
                                                                                                                 AAT17537
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                                                                                                                                                                                                                               20
                                                                                                                                                                                                                                                                                                                      18;
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UNIV UTAH RES FOUND.
US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                              20
                                                                                                                                                                                                                                                                         CAACCTCTGCCTCCCGGGTT 700
                                                                                                                 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 127;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mapping
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                                                                                                                                                                                                                                                                                                                      Conservative
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                            (first entry)
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94US-00308104.
94US-00348824.
95US-00409305.
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95US-00487002.
95US-00488011.
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                                                                                                                 CDNA;
                                                                                                                                                                                                                                                                                                                                         1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                190pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                              3 C; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tavtigian SV,
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                                                                                                                                                                                                                                                                                                                                                                                                                 G; 3
                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                            Score 16.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                 T; 0
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Wiseman RW,
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                                                                                                                                                                                                                                                                                                                                                                   ВB
                                                                                                                                                                                                                                                                                                                                                                                                                 Other;
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Futreal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       partic. breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mapping;
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                                                                                                                                                                                                                                                                                                                         Gaps
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20

CAACCTCTGCCTTCCAGGTT

681 CAACCTCTGCCTCCCGGGTT 700

18;

Conservative

<u>,,</u>

ed. No. 1.6e+03 Mismatches

Indels

0,

Gaps

0

Similarity

RESULT 1283 AAT32608/c

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Query Match
Best Local S
Matches 18
                                                                                                                  can be used as immunogens for antibody production. The mutant BRCA1 genes have at least 1 mutation or polymorphism in comparison to the wild type BRCA1 sequence. By detecting a germline alteration in the BRCA1 gene, a predisposition for breast and ovarian cancer can be diagnosed. In one method, BRCA1 mRNA isolated from a tissue sample from a subject has a probe, corresponding to a fragment of the BRCA1 sequence (or an allelespecific probe for a mutation of it), added to it. The conditions allow for hybridisation of the probe to the mRNA, and any hybridisation which occurs is detected. Alternatively the BRCA1 gene in the tissue sample is isolated, and a shift in electrophoretic mobility of single stranded DNA from the sample on a non-denaturing polyacrylamide gel indicates a mutation. These methods of detection can also diagnose a lesion neoplasia associated with the BRCA1 locus. The methods may be used in gene therapy, protein replacement therapy and protein mimetics, and may be used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-MAR-1995;
07-JUN-1995;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cancer therapy; breast and ovarian cancer predisposing gene; immunogen; antibody production; germline alteration; probe; lesion neoplasia; huma gene therapy; protein replacement therapy; protein mimetic; BRCA1; PCR; polymerase chain reaction; primer; amplify; tandem repeat; 88.
                                                                                                                                                                                                                                                                                                                                                                   AAT17531-T17538 represent amplification primers for tandem repeat markers in the cDNA of the human breast and ovarian cancer predisposing gene (BRCA1) (see AAT17438 for cDNA sequence, and AAT17530 for genomic sequence). This sequence is used in conjuncture with AAT17538 to amplify the short tandem repeat tdj1239. These primers were used in mapping the BRCA1 gene, and for isolating mutations in it. Proteins encoded by mutations of the BRCA1 sequence (see AAT17439-T17453 and AAT17455-T17529) mutations of the BRCA1 sequence (see AAT17439-T17453 and AAT17455-T17529)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-SEP-1994;
16-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer #1 for tandem repeat marker tdj1239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New nucleic acid and polypeptide for mutant or polymorphic BRCA1 for diagnosis and therapy of human breast and ovarian cancer and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-139702/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shattuck-Eidens
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                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnosing pre-disposition to these cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MYRI-)
(RECH-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-NOV-1994;
                                                                                                       screen for drugs in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MYRIAD GENETICS INC.
CENT RECH DU CHUL.
CANCER INST.
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                                                                     BP;
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95US-00409305.
95US-00480784.
95US-00483553.
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94US-00308104
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                                                                     σ
                                                                     P,
                                                                                                       cancer therapy
                1.7%;
                                                                     3 C; 8 G; 3 T;
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                  Score 16.8;
Pred. No. 1
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                                                                       0 Other;
                                      DB 1;
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                                    Length
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for
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RESULT 1284
AAT89640
ID AAT8964
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AC AAT8964
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16-SEP-1994;
29-NOV-1994;
24-MAR-1995;
07-JUN-1995;
07-JUN-1995;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                           The BRCA1 breast/ovarian cancer susceptibility gene has been localised to chromosome 17q. 4 kindred families have provided enough genetic evidence to a sufficiently small region for the application of positional cloning strategies. The primers AAT32602-9 were used to generate a refined physical map of the region surrounding the BRCA1 gene. Esp. the primers AAT32602-3 amplify marker D8178754, AAT32604-5 amplify marker D8178975, AAT32606-7 amplify marker tdj1474 and AAT32608-9 amplify marker tdj1239. The results of the map show that the BRCA1 gene lies between the markers tdj1474 and U5R, an estimated distance of 650 kb. Isolation of the BRCA1 gene, (AAT32601) has allowed development of methods to diagnose a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BRCA1; breast; ovary; cancer; susceptibility; chromosome primer; PCR; amplification; polymerase chain reaction; ge diagnosis; predisposition; ss.
                     AAT89640
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New method for diagnosing a predisposition to breast and ovarian cancer by detecting a germline alteration in the BRCA1 gene or gene regulatory sequence; for gene therapy and to screen for drugs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BRCA1 gene mapping primer tdj1239 A for locus tdj1239.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT32608 standard;
                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 6 A; 3 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 6; Page 127; 190pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Skolnick MH, Goldgar DE, Miki Y, Shattuck-Eidens DM, Tavtigian SV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MYRI-)
(UTAH )
(USSH )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9605308-A1
                                                           AAT89640 standard;
                                                                                                                                                                                                                                                                                                                                                 predisposition to breast and ovarian cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-AUG-1994;
                                                                                                                                                                                             681 CAACCTCTGCCTCCCGGGTT
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                                                                                                                                                                                                                                   18;
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UNIV UTAH RES FOUND.
US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                         CAACCTCTGCCTTCCAGGTT 1
                                                                                                                                                                                                                                   Conservative
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94US-00300266
94US-00308104
94US-00348824
95US-00409305
95US-004093505
95US-004870022
95US-00488011
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                 1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
                                                           20
                                                           BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                   Score 16.8;
Pred. No. 1.
                                                                                                                                                                                             700
                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Swenson
Wiseman
                                                                                                                                                                                                                                                   1.6e+03;
                                                                                                                                                                                                                                                                     DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ŗ, G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         reaction; genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kamb A,
Futreal
                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Harshman
PA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17q; mapping;
enetic marker;
                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ĕ
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 EXSXEXEXEXXXXX
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RESULT 1285
AAV07752
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        p120 is a 120 kD nucleolar antigen protein that is associated with cell proliferation and growth. AAR99630-41 are oligonucleotides antisense to regions of the human p120 gene, that were created and tested for the ability to inhibit the production of p120 and tumour cell growth. These oligonucleotides failed to inhibit p120 production but some did have an effect on tumour cell growth (Hela 33 cells). Other oligonucleotides antisense to the 3'UTR of the p120 gene (see AAT89628 and AAT89629) did inhibit p120 production and the growth of tumour cells in vitro and in vivo. These may be used to treat malignancies, especially human breast cell carcinoma, human epithelioid cervix carcinoma, human amelanotic melanoma and human renal cell carcinoma, and other hyperproliferative of the seasons are informative and cardinoma, and other hyperproliferative of the seasons are first arms of cardinoma, and other hyperproliferative of the seasons.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense; p120; proliferation associated protein; inhibition; UTR; untranslated region; hyperproliferative; cancer; neoplasia; tumour; malignant; carcinoma; melanoma; cardiovascular; inflammation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antisense oligonucleotide specific for pl20 gene 3'UTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-MAR-2003
22-JAN-1998
Synthetic
                                                   phosphorothioate;
antisense; EDITH;
                                                                                                                                                                                    07-DEC-1998
                                                                                                                                                                                                                                     AAV07752;
                                                                                                                                                                                                                                                                                        AAV07752 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diseases, e.g. inflammatory and cardiovascular diseases. (Updated on MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and other hyper-proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anti:sense oligo:nucleotide(s) specific and other hyper-proliferative diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1997-414659/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-FEB-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US5656743-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                              Phosphorothioate oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BAYU ) BAYLOR COLLEGE MEDICINE (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18;
                                                                                                                                                                                                                                                                                                                                                                                                                    1 TCTCGAACACCTGACCTCAG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ļ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TCTCGAACTCCCGACCTCAG 234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 BP; 5 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Col 17; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Saijo Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    92US-00841660.
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                                                                              sulphurisation; heterocycle;
                                                     Beaucage reagent; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Busch RK,
                                                                                                                                                                                                                                                                                        ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 16.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.6e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               p120 -
                                                                              automated synthesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for therapy of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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Key

Location/Qualifiers

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AAV23982/c
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                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention provides a method for sulphurising phosphorus-containing compounds. It comprises contacting the phosphorus-containing compound which a 1,2,4-dithiazolidine-2,5-dione compound or a 3-substituted-1,2,4-dithiazolin-5-one compound. The method is especially useful for incorporation of phosphorothioate linkages into biologically important molecules such as DNA, RNA and phosphopeptides. Molecules containing such linkages are useful e.g. as antisense compounds for inhibiting gene expression, as reagents for studying DNA-protein or RNA-protein interactions, or as catalytic RNA. The present sequence represents an oligonucleotide with phosphorothioate linkages prepared by the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sulphurisation of phosphorus-containing oligo:nucleotide(s) - by contacting the containing five-membered heterocycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9741130-A2
                                                                                                                                         PCR primer; AGT; angiotensinogen; molecular variant detection; essential hypertension predisposition; plasma AGT; G-6A mutation; pregnancy induced hypertension; growth hormone; ss.
                                                                                                                                                        PCR primer;
essential h
                                                                                                                                                                                             Primer for human
                                                                                                                                                                                                                      04-AUG-1998
                                                                                                                                                                                                                                              AAV23982;
                                                                                                                                                                                                                                                                    AAV23982 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 7; Page 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1997-549671/50
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(LOUU ) UNIV LOUISIANA STATE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-APR-1997;
                                                                                                                       Synthetic
                                   07-OCT-1994;
                                                                                   US5763168-A.
            30-SEP-1992;
                                                            09-JUN-1998
                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                          427
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1; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 1 A; 0 C; 0 G; 0 T; 19 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                 1.7%;
larity 5.0%;
Conservative 1
                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     96US-00641920
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            9208-00952442
                                   94US-00319545
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/*tag= a
/note= "phosphorothioate internucleotide linkages"
                                                                                                                                                                                            growth hormone fragment.
                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        51pp; English.
                                                                                                                                                                                                                                                                     20
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                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. 1.6e
17; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  & AGRIC.
                                                                                                                                                                                                                                                                                                                                                                                            Score 16.8;
red. No. 1.6
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                                                                                                                                                                                                                                                                                                                                                                             .6e+03;
2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             compounds, e.g. compound with a dissulphide-
                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ŗ
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                                                                                                                                                                                                                                                                                                                                                                                   Indels
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     PI PRESENTATION OF THE PRE
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AAV85807/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              determination of the predisposition of a human to essential hypertension or pregnancy induced hypertension and comprises, analysing the DNA sequence of the angiotensinogen (AGT) gene for the G-6A mutation, where the presence of the mutation is indicative of a predisposition to essential or pregnancy induced hypertension. The method is useful for the molecular identification of hypertension. The mutation in the AGT gene at position -6 leads to increased plasma AGT concentrations, giving the physiological symptoms for this disease. The mutation (G to A) can be screened for using sequencing methods or hybridisation with a mutation specific primer. Previous disposition to the condition relied on inheritance analysis (ratios, calculations, etc.) between parents/siblings to determine linkage. With the method, a specific diagnosis can be made
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IRP5; LDL-receptor related protein; TRP-3; IDDM; diagnosis; endocytosis;
insulin dependent diabetes mellitus; autoimmune disease;
glomerulonephritis; inflammation; viral infection; osteoporosis;
hypercholesterolemia; Alzheimer's disease; low density lipoprotein;
PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence represents a PCR primer for be used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3; Col 13; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kotelevtsev Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 8 C; 5 G; 3 T; 0 U; 0 Other:
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Todd JA, Hess JW, Co
Hey P, Kawaguchi Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV85807 standard; DNA; 20
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                                                                                  (WELL )
                                                                                                                                                              15-APR-1997;
05-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                      WO9846743-A1
                                                                                                                                                                                                                                                                                                                                                                                                            Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-FEB-1999
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                                                                                                                                                                                                                                             15-APR-1998;
                                                                                                                                                                                                                                                                                                22-OCT-1998.
                                                                                                                                                                                                                                                                                                                                                                                                         sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                  WELLCOME TRUST LTD MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer 58-12 1f.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                 97US-0043553P
97US-0048740P
                                                                                                                                                                                                                                                98WO-GB001102
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  Caskey CT,
Merriman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer for human grwoth hormone, that
he invention. The method is for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 16.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 1.6e+03
     Cox RD, Gerhold TR, Metzker ML,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
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     D, Hammo
Nakagawa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1
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           Hammond
agawa Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              detecting
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AAV85885/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes LRP5 (low density lipoprotein (LDL) receptor related protein, previously designated LRP-3). AAV85827 to AAV85822 represent exon primers used for obtaining LRP5 cDNA. Nucleic acid molecules (NAMS) encoding LRP5 can be used for determining if an individual is susceptible to insulin dependent diabetes mellitus (IDDM). The NAMS or proteins can be used for reducing triglyceride levels in the serum of an individual. Therapies that affect LRP5 may also be useful in the treatment of autoimmune diseases such as glomerulonephritis, diseases and disorders involving disruption of endocytosis and/or antigen presentation, cytokine clearance and/or inflammation, viral infection, pathogenic bacterial toxin contamination, elevation of free fatty acids or hypercholesterolemia, type 2 diabetes, osteoporosis, Allheimer's disease and cardiovasgular disease. Products from the present invention
                                                                                                                                                                                                                                                                                                                                              LRP5; LDL-receptor related protein; LRP-3; IDDM; diagnosis; endocytosis; insulin dependent diabetes mellitus; autoimmune disease; glomerulonephritis; inflammation; viral infection; osteoporosis; hypercholesterolemia; Alzheimer's disease; low density lipoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated LDL-receptor related protein - used to develop products treating, e.g. elevated triglyceride levels, diabetes, autoimmune disorders, inflammation or Alzheimer's disease.
                                        Todd JA, Hess JW, Caske
Hey P, Kawaguchi Y, Mer
Phillips MS, Twells RCJ;
                                                                                                                                                                                                                                                                                                                                                                                                                           LRP5 SNP primer 58-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV85885 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 5 A; 6 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 12; Page 106; 200pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Phillips MS,
            WPI; 1998-594573/50
                                                                                                      (WELL )
                                                                                                                                                                                                                                                           W09846743-A1
                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV85885;
                                                                                                                                                   15-APR-1997;
05-JUN-1997;
                                                                                                                                                                                                                             22-OCT-1998
                                                                                                                                                                                              15-APR-1998;
                                                                                                                                                                                                                                                                                         Homo
                                                                                                                                                                                                                                                                                                                                   primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 also be used for detection, diagnosis and drug screening
                                                                                                                                                                                                                                                                                       sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 AGCGGTGTGATCTCAGCTCA 1
                                                                                                      MERCK & CO
                                                                                                       WELLCOME TRUST LTD
MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGTGGTGTGATCACAGCTCA 503
                                                                                                                                                                                                                                                                                                                                     88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                  97US-0043553P
97US-0048740P
                                                                                                                                                                                               98WO-GB001102
                                                                     Caskey CT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.7%;
                                                           Merriman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ₽₽
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 16.8; DB 1
Pred. No. 1.6e+03
                                                        TR,
                                                        RD, Gerhold D, Hammo
Metzker ML, Nakagawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
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                                                          Hammond
agawa Y;
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New isolated LDL-receptor related protein - used to develop products treating, e.g. elevated triglyceride levels, diabetes, autoimmune disorders, inflammation or Alzheimer's disease.
                                                                                                                                          Claim 12; Page 111; 200pp; English.
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the treatment of autoimmune diseases such as glomerulonephritis, diseases and disorders involving disruption of endocytrosis and/or antigen presentation, cytokine clearance and/or inflammation, viral infection, pathogenic bacterial toxin contamination, elevation of free fatty acids or hypercholesterolemia, type 2 diabetes, osteoporosis, Alzheimer's disease and cardiovascular disease. Products from the present invention The present invention describes LRP5 (low density lipoprotein (LDL) receptor related protein, previously designated LRP-3). AAV85823 to AAV85900 represent SNP primers used for obtaining LRP5 CDNA. Nucleic acid molecules (NAMs) encoding LRP5 can be used for determining if an individual is susceptible to insulin dependent diabetes mellitus (IDDM). The NAMs or proteins can be used for reducing triglyceride levels in the serum of an individual. Therapies that affect LRP5 may also be useful in Sequence can also 20 BP; 5 be used for detection, diagnosis and drug screening A; 6 C; 5 G; 4 T; 0 U; 0 Other; Nucleic acid

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Query Match
Best Local S
Matches 18
                484 AGTGGTGTGATCACAGCTCA 503
20
                                  18;
                                          Similarity
AGCGGTGTGATCTCAGCTCA 1
                                   Conservative
                                          1.7%;
                                 0;
                                          Score 16.8;
Pred. No. 1.
                                  Mismatches
                                          .6e+03
                                                    DB 1;
                                   <u>.</u>
                                                   Length 20;
                                   Indels
                                   0
                                 Gaps
                                   0
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RESULT 1289
AAV09200
Type I interleukin-1 receptor; IL1R; human; IL1 protein; hybridisation; inflammation; ss; 3' untranslated region; phosphorothioate linkage.
                                                                                                                                                       09-JUN-1998
                                                                                           Phosphorothiate oligonucleotide sequence 8802 targeting ILIR mRNA.
                                                                                                                                                                                                                    AAV09200;
                                                                                                                                                                                                                                                                             AAV09200 standard; DNA;
                                                                                                                                                       (first entry)
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Key
modified_base
                                               WO9744656-A1.
                                                                Homo
                                                                sapiens.
                                                           Location/Qualifiers
                                                    /note= "Phosphorothioate internucleotide linkage"
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## 27-NOV-1997

97WO-US007147. 96US-00651692. PHARM INC. Bennett CF, Dean N, Geiger T;	Miraglia L,	(ISIS-) ISIS PHARM INC.	21-MAY-1996;	12-MAY-1997;
	Dean N, Geiger	PHARM INC.	96US-00651692.	97WO-US007147.

WPI; 1998-018646/02.

2'-substituted oligonucleotide(s) specific for interleukin-1 receptor type I - used to modulate expression and detect overexpression of the Example 5; receptor. Page 19; 63pp; English

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RESULT 1290
AAZ37721/c
ID AAZ37721/c
XX AAZ37722
XX AAZ3772
XX AAZ3772
XX Human m
XX (ISIS-)
XX (ISIS-)
XX Hiragli
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XX Hiragli
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This is a novel oligomer comprising 20 covalently linked nucleotides which bind to the 3' untranslated region of the interleukin-1 receptor (IIIR) mRNA. Expression of IIIR, in cells and tissues can be modulated be compositions comprising oligomers which are able to specifically hybridise with target areas of its encoding sequence. The composition cape used for treatment of disease in humans caused by excessive receptor expression, e.g. inflammation. When labelled they can be used diagnostically to determine overexpression of IIIR, also to determine localisation and distribution of this expression for research, diagnostic
                                                    AAZ37473-Z37738 represent human mdm2 phosphorothioate oligonucleotides. AAZ37471, AAZ37472, AAZ37739, AAZ37740 and AAZ37741 are used in the exemplification of the present invention. The present invention describes novel nucleotide antisense compounds, targetted to the 5' untranslated, translation termination codon, or 3' untranslated region of a nucleic acid encoding human mdm2, that modulates expression of human mdm2. The oligonucleotides mediate their effect by antisense inhibition of hyperproliferative gene expression. The antisense compound is used to treat an animal having a disease or condition associated with mdm2, particularly a hyperproliferative condition, more particularly cancer, especially of the blood, brain, breast, lung or soft tissue, or psoriasis, fibrosis, atherosclerosis or restenosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antisense;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JAN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            restenosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      lung cancer; soft tissue cancer; psoriasis; fibrosis; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapeutic purposes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                            54; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                compounds used to treat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nero
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Pred. No. 1.6e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hyperproliferative conditions
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Sequence

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6 A;

3 C;

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4 T;

0 U;

0 Other;

RESULT 1292 AAZ37735/c

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RESULT 1291
AAZ37728/c
ID AAZ37728/c
ID AAZ3772
XX AAZ3772
XX AAZ3772
XX Human n
KW Human n
KW Human n
KW Ilung cc
KW restenc
XX Synthet
OS Homo sc
XX W099490
XX W099490
XX (ISIS-)
XX (ISIS-)
XX Miragl:
XX Miragl:
XX New ant
XX PR Exampl
CC AAZ374
CC AAZ374
CC CAAZ374
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                           316 GTAGAAACAGGGTTTCACTG
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expression; inhibition;
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Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human mdm2 gene; proliferation; tumour; phosphorochioate; p53; cancer; antisense; modulation; oligonucleotide; expression; inhibition; hyperproliferation; blood cancer; brain cancer; breast cancer; ancer; sense cancer; brain cancer; streast cancer; brain cancer; streast cancer; brain cancer; soft tissue cancer; psoriasis; fibrosis; atherosclerosis;
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Human mdm2 gene; proliferation; tumour; phosphorothicate; p53; cancer; antisense; modulation; oligonucleotide; expression; inhibition; hyperproliferation; blood cancer; brain cancer; breast cancer; hyperproliferation; blood cancer; psoriasis; fibrosis; atherosclerosis;
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                                                                                                                                                                  androgenetic alopecia; alopecia areata; alopecia universalis; wildtype;
hair follicle; hairless; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This primer can be used in the specific PCR amplification of the human CC hairless intron 15. This PCR allowed the sequencing of intron 15 and CC comparision of the nucleotide sequence. A mutation was found within this cintron that after further analysis was associated with the alopecia CC universalis phenotype in this family. The gene was discovered by CC genotyping a Pakistani kindred (comprising of 4 affected males and 7 CC inheritance. The invention provides methods and sequences for the CC recombinant production of wild-type human hairless, mutant human hairless and wild-type human whn (winged-helix-nude) proteins, assays for CC and wild-type human whn (winged-helix-nude) proteins, assays for CC screening for binding compounds, modulators and homologues, and animal CC alopecia (male pattern baldness), alopecia areata, alopecia totalis, CC congenital alopecia universalis, congenital alopecia and severe T-cell immunodeficiency can be treated with compounds identified in the assays. The methods are also useful for identifying compounds that can be used to inhight hair process.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1295
AAV80037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Christiano AM
                                                                                                                                                                                                                                                                                        Phosphomannomutase-2; PMM2; CDG1; mutation; human; transgenic; assay; carbohydrate-deficient glycoprotein syndrome type 1; drug screening; Jaeken disease; single-strand confirmation polymorphism; SSCP;
                                                                                                                                                                                                                                                                                                                                                                                  16-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV80037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inhibit hair growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 42; 127pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hair growth.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human hairless gene and
                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                          prenatal diagnosis; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                     Primer int4R for
             WPI; 1999-024063/02
                                                                                                   30-APR-1997;
27-JAN-1998;
                                                                                                                                              30-APR-1998;
                                                                                                                                                                        05-NOV-1998
                                                                                                                                                                                                     WO9849324-A2
                                          Matthijs
                                                                        (GENZ ) GENZYME
                                                                                                                                                                                                                                 sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               391
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                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGTGCCAGGATTACAGGCGT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGTGCTGGGATTACAGGCGT 410
                                          G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                  .(first entry)
                                                                                                   97GB-00008851.
98GB-00001719.
                                                                                                                                              98WO-EP002593.
                                                                        UK LTD
                                                                                                                                                                                                                                                                                                                                                     SSCP analysis of PMM2 exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein, useful for identifying modulators of
                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>.</u>.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.8; DB 1;
Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                     58
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New DNA encoding human phosphomannomutase or its fragments - used to detect mutations associated with carbohydrate-deficient glycoprotein syndrome-1, particularly for prenatal diagnosis.
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Claim 5; Page 64; 104pp; English

express PMM2 or its mutants; and to create transgenic animals for use in drug screening and for studying expression pathways. The expressed proteins are used to screen for agents that modulate activity of PMM2, proteins are used to screen for agents that modulate activity of PMM2, for therapy and to raise specific antibodies (for detecting PMM2 or its mutants, in competitive or capture assays). Biochemical assays for phophomannomutase activity are used to identify possible carriers of CDG1 (Jacken disease). Measuring enzymatic activity in foetal cells (or in parental leucocytes if such cells are unavailable) and detecting mutations in the PMM2 gene makes possible a better prenatal diagnosis of CDG1. Sequences AAV80026-43 represent primers used in PCR and singlestrand confirmation polymorphism (SSCP) analysis of the 8 exons of PMM2 gene. These primers are used to determine the SSCP mutations in the PMM2 are sequences invention relates to a human phosphomannomutase-2 (PMM2) protein and nucleoride sequence encoding the protein. The DNA or its fragments used to detect mutation in the PMM2 genes that are associated with carbohydrate-deficient glycoprotein syndrome type 1 (CDG1). The usences can also be used to detect expression of PMM2-related cDNA; to ii 6

Sequence 20 BP; 4 A; 2 C; 8 G; 6 T; 0 U; 0 Other;

Matches Query Match Best Local : 392 GTGCTGGGATTACAGGCGTG 411 1 Similarity 18; Conserv Conservative 1.7%; <u>.</u> Score 16.8; Mismatches No. 1.6e+03 DB 1; Length Indels 20; 0, Gaps 0

RESULT 1296

AAV80023 standard; DNA;

20

밁 Ś

GTGTTGGGATTACAGGCATG 20

AAV80023;

16-MAR-1999 (first entry)

Exonic primer PMM16-int5R for PMM2 SSCP analysis.

Phosphomannomutase-2; PMM2; CDG1; mutation; human; transgenic; assay; carbohydrate-deficient glycoprotein syndrome type 1; drug screening; Jaeken disease; single-strand confirmation polymorphism; SSCP; prenatal diagnosis; PCR primer; ss.

Homo Synthetic sapiens

WO9849324-A2

05-NOV-1998.

30-APR-1998; 98WO-EP002593

30-APR-1997; 27-JAN-1998; 97GB-00008851. 98GB-00001719.

(GENZ ) GENZYME UK LTD

Matthijs G;

WPI; 1999-024063/02.

New DNA encoding human phosphomannomutase or its fragments - detect mutations associated with carbohydrate-deficient glyco syndrome-1, particularly for prenatal diagnosis nts - used to glycoprotein

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RESULT 1297
AAZ18580/c
ID AAZ18580/c
ID AAZ1858
AC AAZ1811
AC AAZ1811
CC The in
CC The in
CC AAZ1811
CC AAZ181
CC AAZ1811
CC AAZ1811
CC AAZ1811
CC AAZ1811
CC AAZ1858

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a human phosphomannomutase-2 (PMM2) protein and the nucleotide sequence encoding the protein. The DNA or its fragments are used to detect mutation in the PMM2 genes that are associated with the carbohydrate-deficient glycoprotein syndrome type 1 (CDG1). The sequences can also be used to detect expression of PMM2-related cDNA; to express PMM2 or its mutants; and to create transgenic animals for use in drug screening and for studying expression pathways. The expressed proteins are used to screen for agents that modulate activity of PMM2, for therapy and to raise specific antibodies (for detecting PMM2 or its mutants, in competitive or capture assays). Biochemical assays for phophomannomutase activity are used to identify possible carriers of CDG1 (Jaeken disease). Measuring enzymatic activity in foetal cells (or in parental leucocytes if such cells are unavailable) and detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CDG1. The present sequence represents an exonic primer used for the single-strand confirmation polymorphism (SSCP) analysis of PMM2 exon
                                                                                                                                                                                                                                      Disclosure; Page 50; 195pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                            Brooks-Wilson AR,
Miller A, North M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9937809-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ASTH1; asthma; human; chromosome 11p; ASTH1I; ASTHIJ; genetic locus; ss; therapeutic; immunogen; polymorphism; PCR primer; microsatellite marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ18580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ18580 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mutations in the PMM2 gene makes possible a better prenatal diagnosis of
                                                                                                                                                                                                                                                                                                                        Mammalian asthma related genes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (AXYS-) AXYS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JUL-1999
                                                                                                                                                                                                                                                                                                development of asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          392 GTGCTGGGATTACAGGCGTG 411
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ASTH1 polmorphic microsatellite marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 4 A; 2 C; 8 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 14; 104pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98WO-US001260
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Buckler A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cardon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                           useful for diagnosis of a predisposition
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Carey
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Galvin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 88;
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The invention identifies a genetic locus ASTH1, associated with asthma, mapped to human chromosome 11p. ASTH11 and ASTH1J are genes present within the locus, located close to each other on human chromosome 11p, and have similar patterns of expression, and common sequence motifs. The ASTH1 genes and fragments, encoded protein, genomic regulatory regions and anti-ASTH1 antibodies are useful in the identification of individuals

predisposed to development of asthma,

and for

modulation of

RESULT 1299 AAA96399/c ID AAA9639

AAA96399 standard; DNA;

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ВP

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RESULT 1298
AAZ43583/c
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밁
                                                   Query Match
Best Local S
Matches 18
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Best Local
                                                                                                                                         This invention describes a novel method for determining the danger of suffering from Alzheimer's disease (AD) in which if the 19117th to the 19183rd bases in the gene of a dihydrolipoamidosuccinyl transferase of a mitochondria alpha-ketoglutarate dehydrogenase complex are respectively A and C in the above order in both genes derived from its father and mother is checked. The method is useful for the prevention of Alzheimer's disease. AAZ435/4-Z43603 represent primers used in the detection method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   activity in vivo for prophylactic and therapeutic purposes. The ASTI protein is useful as an immunogen to raise specific antibodies, in screening for compositions that mimic or modulate ASTH1 activity or expression, including altered forms of ASTH1 protein, and as a therapeutic. Sequences AAZ18510-Z18631 represent PCR primers for polymorphic microsatellite markers in the ASTH1 region
                                                                                                                                                                                                                                                                              Determination of danger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 6 A; 3 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                      WPI; 2000-046934/04.
                                                                                                                                                                                                                                                                                                                                                                              28-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                       28-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                 09-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                          JP11308996-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mitochondria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Alzheimer's disease; primer; dihydrolipoamidosuccinyl transferase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alzheimer's disease detecting primer #10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ43583;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ43583
                                                                                                                                  described in the invention
                                                                                                                                                                                                                                                       Example;
                                                                                                                                                                                                                                                                                                                                                   (SRLS-) SRL KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             931 CTCACTCTGTTACCCAGGCT 950
                         314
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                                                    18;
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                      TGGTAGAAACAGGGTTTCAC 333
                                                                                                     20 BP; 5 A;
                                                                                                                                                                                                                                                      Page 7; 9pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTCACTCTGTCTCCCAGGCT 1
 TAGTAGAGACAGGGTTTCAC 1
                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              alpha-ketoglutarate dehydrogenase; detection; ss.
                                                                                                                                                                                                                                                                                                                                                                              98JP-00134578
                                                                                                                                                                                                                                                                                                                                                                                                       98JP-00134578
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.7%;
                                                                90.0%;
                                                                                                       6 C; 3
                                                                             1.7%;
                                                                                                                                                                                                                                                                               of suffering from Alzheimer's disease in genes encoding enzyme derived from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>.</u>
                                                                                                       G; 6 T; 0 U; 0
                                                Pred. No. 1.66
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16.8;
Pred. No. 1.
                                                                  Score 16.8;
Pred. No. 1
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                                                                  .6e+03
                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                         Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                             Length
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                                                      Indels
                                                                               20;
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                                                                                                                                                                                                                                                                                  parents.
                                                    Gaps
                                                                                                                                                                                                                                                                                             comprises
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RESULT 1300
AAZ43613
ID AAZ4361
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AC AAZ4361
XX
AC AAZ4361
XX
DT 22-FEB-
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                       CC microsatellite repeat (PMR) sequences from the human costimulatory creeptor gene locus (hCGRL). The primers are used in the method of the cinvention. The specification describes a method for determining the creeptor gene locus (hCGRL). The primers are used in the method of the comprises detecting a pMR sequence in the CD28, ICOS gene or CTLA4 (C gene of the human costimulatory receptor gene locus (hCGRL). PMR CC generate products that differ in size. These products can then be compared by rapid and convenient high resolution processes. The method is useful for determining the predisposition of insulin-dependent diabetes mellitus (IDDM), Addison's disease, Graves disease, autoimmune chipothyroidism, myasthenia gravis, thymoma, lugus, thyroiditis, coeliac disease and leprosy. PMR sequences within hCRGL are useful as coeliac disease and leprosy. PMR sequences within hCRGL are useful as coeliac disease and leprosy. PMR sequences within hCRGL are useful as coeliac disease and leprosy. PMR sequences within hCRGL are useful as coeliac disease.
                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA96399;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primers AAA96399-A96400 were used to amplify polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 18; Page 152; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Determining predisposition of humans to develop autoimmune disease involves detecting polymorphic microsatellite repeat sequence within
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-628257/60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hashimoto's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primer used to amplify a sara31/32 polymorphic microsatellite repeat.
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                        22-FEB-2000
                                                                  AAZ43613;
                                                                                                         AAZ43613
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                                                                                                                                                                                                                                                                                            18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                          diagnosis and human
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                                                                                                                                                                                                                                              GTCACCCAGGCTGGAGTGCA 658
                                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                                         standard;
                                                                                                                                                                                                               GTCGCTCAGGCTGGAGTGCA 1
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                                                                                                                                                                                                                                                                                                                                                                                  BP;
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                        (first entry)
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                                                                                                         DNA;
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                                                                                                                                                                                                                                                                                                              1.7%;
                                                                                                                                                                                                                                                                                                                                                                                  8 C; 5
                                                                                                            20
                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                        genome mapping
                                                                                                                                                                                                                                                                                                                                                                                  G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                  Score 16.8;
Pred. No. 1.
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06-JAN-2000.

WO200000647-A1 Homo sapiens.

15-APR-1999;

99WO-US008280

Human; angiotensinogen gene; hypertension; upstream stability factor; oestrogen recept

receptor; GH gene; PCR primer;

88.

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transcription factor;

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RESULT 1301
AAZ60319/c
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Best Local :
                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes a novel method for detecting the presence in a subject of a polymorphism linked to a gene associated with familial dysauconomia comprising analyzing human chromosome 9. The method comprises analyzing human chromosome 9 for the presence of a polymorphism located between D9S3 and D9S105 inclusive and linked to the gene associated with familial dysautonomia where the presence of a polymorphism is indicative of carriers of a gene associated with familial dysautonomia. The methods allow characterization of simple sequence repeat polymorphisms using less DNA, typically 10 nanograms of genomic DNA, and is faster than restriction fragment length polymorphism analysis. AAZ43609-Z43642 represent PCR primers used in the detection method described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detection; polymorphism; familial dysautonomia; human; chromosome p9S53; D9S105; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Breakefield
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16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human familial dysautonomia D9S58 marker PCR primer
PCR primer used
                                                                                                                              AAZ60319 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Col
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting polymorphisms linked to a gene associated with familial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-052539/04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                          05-MAY-2000
                                                                                     AAZ60319;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                  725 CCTGAGTAGCTGGGACTACA 744
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                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                          CCTGAGTAGCCGGGACTATA 20
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                                                                                                                                                                                                                                                                                                                                                                                                                BP; 5
                                                                                                                                                                                                                                                                                                                               Conservative
                                          (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     92US-00890719.
93US-00049678.
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to amplify part of the human GH gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Slaugenhaupt S,
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                                                                                                                                                                                                                                                                                                                                                                                                                  C; 6 G; 4 T; 0 U;
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                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                   Score 16.8;
Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                     0 Other;
                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                          Length
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                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                 Gaps
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RESULT 1302
AAA54272/c
ID AAA5427
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primers AAZ60319-20 were used to amplify part of the human GH (not specified) gene. The specification describes a method for the prognosis of a known predisposition to hypertension, which is associated with the angittensinogen gene in humans and is caused by the presence of a mutation at -20 from C to A (A-20C mutation). Mutation at this position reduces affinity for the following transcription factors: upstream stability factor and oestrogen receptor. The presence of this mutation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in a subject predisposed to hypertension, indicates increased predisposition, or predisposition to more severe disease. The method is allows affected subjects to be monitored closely, and treated, before the second second subjects to be monitored closely.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-JUN-1998;
                                                  New antisense compounds that hybridizes with and inhibits the of human telomeric repeat binding factor 1 (TRBF-1), useful for conditions or diseases associated with TRBF-1 expression.
                                                                                                                                                                                                                                                                                                                                               Human telomeric repeat binding factor 1; hTRBF-1; antisense; disease; modulation; expression; prophylaxis; infection; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20
                          Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA54272;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA54272 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prognosis of hypertension, associated with the angiotensinogen gene,
                                                                                                                                      Monia BP,
                                                                                                                                                                                           21-JUL-1999;
                                                                                                                                                                                                                       21-JUL-1999;
                                                                                                                                                                                                                                                  10-OCT-2000.
                                                                                                                                                                                                                                                                                                                                  anti-inflammatory; tumour;
                                                                                                                                                                                                                                                                                                                                                                                         Antisense
                                                                                                                                                                                                                                                                                                                                                                                                                    26-FEB-2001
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                                                                                                           2000-664192/64.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                        Col 63; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTCCGAGGCTGGAGTGCAGT 1
                                                                                                                                                                                                                                                                                                                                                                                       oligonucleotide directed against hTRBF-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      becomes serious
                                                                                                                                     Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 4 A; 8 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                subjects at increased risk.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jeunemaitre
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                                                                                                                                                                 PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                    (first
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                                                                                                                                                                                             99US-00358384.
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                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
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                                                                                                                                                                                                                                                                                                                                  diagnostic; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 16.8; DB 1;
Pred. No. 1.6e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
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                                                                 te expression for treating
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Antisense compounds directed against the start codon,

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untranslated

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AGT gene contribute

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RESULT 1303
AAZ38449/c
ID AAZ3844
XX AAZ3844
XX AAZ3844
XX Angiote
KW Angiote
KW Variant
KW detecti
KW linkage
XX Index
OS Synthet
OS Synthet
OS Homo 82
XX Index
ON US59981
XX O7-DEC-
XX O7-DEC-
XX O7-DEC-
XX INRM
PA (INRM)
PR 07-OCT-
XX (UTAH)
PA (INRM)
P
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Best Local Similarity
Matches 18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      its expression. They may also be used prophylactically to prevent or delay infection, inflammation or tumour formation, or as research reagents and diagnostics, e.g. to distinguish between functions of various members of a biological pathway. The antisense oligonucleotides disclosed are described in GENESEQ records AAA54242-A54280, AAA54385
                      This sequence represents hGH-A1819 PCR primer #1, used with primer #2 (AAZ38450) to amplify a portion of the human growth hormone (hGH) gene for linkage analysis with the angiotensin-converting enzyme (ACE) gene. Genetic studies revealed that the genes encoding two key enzymes in the angiotensin II synthetic pathway, renin and ACE, were not associated with human hypertension; however, the angiotensinogen (ACT) gene was involved in the pathogenesis of essential hypertension. Sequence variations in the ACT gene can be identified via amplification of gene fragments via PCR, using primers AAZ3841-Z38448, and subsequent sequence analysis.
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07-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Angiotensinogen; hypertension; pathogenesis; exon; allele; mutation; variant; susceptibility; predisposition; essential; pregnancy-induced; detection; diagnosis; management; ACE; angiotensin-converting enzyme; linkage; analysis; growth hormone; hGH; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ38449 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Analyzing the DNA sequence mutation A-20C is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                      Example 3; Col 13;
                                                                                                                                                                                                                                                                                                                                                                                                                              mutation A-20C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-052541/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lalouel J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UTAH ) UNIV UTAH RES FOUND.
(INRM ) INSERM INST NAT SANT
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94US-00319545.
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                                                                                                                                                                                                                                                                                                                                      25pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                              ce of the angiotensinogen (AGT) gene for determining a predisposition to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Soubrier F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.8; DB 1;
Pred. No. 1.6e+03;
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     RESULT 1304
AAA11942
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene can be used to identify individuals with a genetic predisposition develop essential hypertension or pregnancy-induced hypertension. Detection of a predisposition would then allow specific management of hypertension in these subjects e.g., by dietary sodium restriction, by monitoring blood pressure and treating with conventional drugs, by administration of renin inhibitors or by administration of drugs to inhibit synthesis of AGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       susceptibility to
gene can be used t
develop essential
antiinfectious activity. Synthesized chimeric oligonucleotides targeted to human MDMX, 20 nucleotides in length, composed of a central gap region consisting of ten 2'-deoxynucleotides flanked on both sides by 5-nucleotide wings were tested for antisense inhibition of MDMX expression. Results of real-time quantitative polymerase chain reaction (PCR) showed 71 out of the 159, 20 base pair sequences, all fully defined in the specification, demonstrated at least 30% inhibition of MDMX expression. The antisense oligonucleotides are useful for effective and specific modulation, particularly inhibition of MDMX expression, and may be used in treating humans or animals suspected of having or being prone to a disease or condition associated with expression of MDMX. The antisense oligonucleotides may also be used as research reagents or kits, and as disgnostices, e.g. to elucidate the function of a particular gene or to distinguish between functions of various members of a biological pathway.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 4 A; 8 C; 5 G;
                                                                                                                                                                                                                                                in length, targeted to a nucleic acid encoding a human MDMX. (I) specifically hybridizes with and inhibits the expression of human MDMX. The products of the invention have anticarcinogen, antiinflammatory and
                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides targeting nucleic acids encoding human MDMX useful for inhibiting MDMX expression and for treating diseases associated with MDMX expression e.g. tumor formation, inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antiinfectious;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MDMX; human; antisense; inhibitor; anticarcinogen; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human MDMX antisense oligonucleotide #31065
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-282710/24.
                                                                                                                                                                                                                                                                                                           This invention describes a novel antisense compound (I),
                                                                                                                                                                                                                                                                                                                                               Example 15; Col 97-98; 51pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     modulation; treatment; disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the development of hypertension. Analysis of the AGT
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RESULT 1305
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                                              CC in length, targeted to a nucleic acid encoding a human MDMX. (I) specifically hybridizes with and inhibits the expression of human MDMX. (C) specifically hybridizes with and inhibits the expression of human MDMX. (C) The products of the invention have anticarcinogen, antiinflammatory and contininfectious activity. Synthesized chimeric oligonucleotides targeted to human MDMX, 20 nucleotides in length, composed of a central gap region consisting of ten 2'-deoxynucleotides flanked on both sides by 5-consisting of ten 2'-deoxynucleotides flanked on both sides by 5-consisting of ten 2'-deoxynucleotides flanked on both sides by 5-consisting of ten 2'-deoxynucleotides flanked on both sides by 5-consisting of ten 2'-deoxynucleotides in hibition of MDMX expression. CC results of real-time quantitative polymerase chain reaction (PCR) showed close of the sides of the sides of the sides of the constrated at least 30% inhibition of MDMX expression. The antitions of month and may be used as prophylation of month and specific conditation, particularly inhibition of MDMX expression, and may be used as collagonousleotides may also be used as research reagents or kits, and as diagnostics, e.g. to elucidate the function of a particular gene or to condition of the mathod of the invention of antisense oligonucleotides may also be used as research reagents or kits, and as collagonation. AAA11781-A11945 represent antisense oligonucleotides flanked of the invention of the particular specifical pathway.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense oligonucleotides targeting nucleic acids encoding MDMX useful for inhibiting MDMX expression and for treating disa associated with MDMX expression e.g. tumor formation, inflammati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MDMX; human; antisense; inhibitor; anticarcinogen; antiinflammatory; antiinfectious; modulation; treatment; disease; diagnosis; primer; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human MDMX antisense oligonucleotide #31222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA11941 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2000-282710/24.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Col 97-98; 51pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and for treating diseases formation, inflammation.
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Sequence

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RESULT 1306
AAA80487/c
ID AAA8048
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The invention relates to the ASTH1 locus on the short arm of human CC chromosome (11p). This locus comprises the ASTH1I and ASTH1I genes, which CC are associated with a genetic predisposition to asthma and bronchial CC hyperreactivity. The ASTH1I and ASTH1J genes are oriented in opposite CC directions with the ASTH1 locus, and have similar patterns of expression CC and common sequence motifs. They are both expressed in trachea, lung and Several other tissues. ASTH1I and ASTH1J are novel members of the ets CC family of transcription factors, which have been implicated in the CC genes known to be important in the actiology of asthma. Both ASTH1I and ASTH1J mRNAs are alternatively spliced. Alternative splicing of CC transcripts has no effect on the open reading frame of ASTH1I, as the CC exons involved are all 5' to the start codon in exon b. In contrast, CC alternative splicing of ASTH1I transcripts results in 3 different ASTH1I concluded are all 5' to the start codon in exon b. In contrast, colorisms. The invention also encompasses mouse asth1j protein. The ASTH1I CC predisposition to asthma, as probes for identify a hereditary a hereditary protein in cell lines. The encoded ASTH1 proteins are useful as communogens to raise specific antibodies; in drug screening for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 18; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acids other than naturally occurring chromosomes encoding ASTH1 protein, for e.g. screening compositions that modulate expressior function of ASTH1 proteins or as diagnostics for genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ASTH1 locus; ASTH1I; ASTH1J; human; chromosome 11p; asthma; bronchial hyperreactivity; ets family; transcription factor; splice variant; genetic predisposition; polymorphism; antibodrug screening; prophylaxis; therapy; diagnosis; polymorphic microsatellite marker flanking sequence;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA80487 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Galvin M, Mil
Brooks-Wilson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-JAN-1997;
01-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ASTH1 polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                           Example; Col 31-32; 131pp; English
                                                                                                                                                                                                                                                                                                                                                                                                              predisposition to asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-505109/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US6087485-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (AXYS-) AXYS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        , Miller A, NC-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TCTGTCTCCCAGGCTGAAGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TCTGTCACCCAGGCTGGAGT 655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97US-0035663P.
97US-0051432P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98US-00009913
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    microsatellite marker CA39_2 primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  genotypes; BAGs;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               North M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
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Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cardon L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Buckler
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                             expression
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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RESULT 1307
AAC61094/c
ID AAC6109
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Best Local S
Matches 18
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                                                               cell comprises a recombinant expression construct encoding a mammalian p21 gene which gives rise to the expression of p21 in the cell. The p21 gene is under the control of a promoter. The recombinant cell is used in methods for identifying genes involved in cell cycle progression, growth promotion, modulation of apoptosis, cellular senescence and aging and for identifying compounds that inhibit or potentiate cellular senescence, regulated by p21. The fibrosarcoma cell can be used to produce or an antiapoptotic or mitogenic factor. The present sequence represents a PCR primer used to amplify a human promoter DNA sequence for use in the
                                                                                                                                                                                                                                                                                                                                                                                                                                            Recombinant mammalian fibrosarcoma cell for identifying compounds that inhibit or potentiate cellular senescence, regulated by p21, comprises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAC61094 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         batched analysis of genotypes (BAGs)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-APR-1999;
29-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-APR-2000; 2000WO-US009286.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200061751-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fibrosarcoma cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC61094;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 compositions that mimic or modulate activity or expression of ASTH1I
                                                                                                                                                                                                                                                                                                                                                               Example 6; Page 65; 119pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                      recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-638567/61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer used for amplification of the PLK1 promoter sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-OCT-2000
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                                                                                                                                                                                                                                                                                                                invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   promotion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTCACTCTGTTACCCAGGCT 950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTCACTCTGTCTCCCAGGCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Roninson IB;
                                                                                                                                                                                                                                                                                                                                                                                                                   expression construct encoding a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0128676P
99US-00449589
                                                                                                                                                                                                                                                                                                                relates to a recombinant mammalian fibrosarcoma cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   p21; promoter; cell cycle progression; PCR primer;
apoptosis modulation; senescence; aging; PLK1; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3 C; 9 G; 2 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 16.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                              comprises
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Sequence 20 BP; 4 A; 6 C; 4 G;

6 T; 0 U; 0 Other

expression

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RESULT 1308
AAK95190
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                              RESULT 1309
AAK95165
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Best Local S
Matches 18
                                                                                                                     Query Match
Best Local S
Matches 18
                                                                                                                                                                                              The invention relates to primers for synthesising full length cDNA clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful for clarifying the function of the protein encoded by the cDNA. The full length clones were obtained by construction of full length enriched cDNA libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA easily without any special methods. The present sequence is a primer used to amplify a human cDNA are closed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-JUL-1999;
11-JAN-2000;
                                                                                                                                                                                                                                                                                                                                              830 Primers useful for synthesizing full length cDNA clones
                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-524255/58
                                                                                                                                                                                                                                                                                                                                                                                            Wakamatsu
                                                                                                                                                                                                                                                                                                                                                                                                        Ota T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUL-2000; 2000EP-00114089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human cDNA clone-specific primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAK95190 standard;
                                                                                                                                                                 Sequence
AAK95165
                      AAK95165
                                                                                                                                                                                                                                                                                                              Example 18; Page 133; 1380pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                             (HELI-) HELIX RES INST.
                                                                                                                                                                                                                                                                                                                                     in genetic
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                                                                                                                                Similarity
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                                                                                          CTCAGCCTCCCAAGTAGCTG 561
                      standard;
                                                                            CTCAGCCTTCCAAGTAGCAG 20
                                                                                                                                                                 20 BP; 5
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                                                                                                                                                                                                                                                                                                                                     manipulation.
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                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000JP-00183765
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2000JP-00118774.
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                                                                                                                                                                                                                                                                                                                                                                                           Sugiyama T,
                                                                                                                                                                                       in the invention
                                                                                                                                                                 A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cDNA; cDNA synthesis; oligo-capping; PCR primer; ss.
                      DNA;
                                                                                                                                                                 7 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                     Isogai T,
                                                                                                                                           1.7%;
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                      BP.
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Pred. No. 1.6e+03;
                                                                                                                                Score 16.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                            Hayashi K,
K, Kojima
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                                                                                                                                 1.6e+03
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S, Otsuki
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T, Koga
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RESULT 1310
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11-JAN-2000;
02-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to primers for synthesising full length cDNA clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful for clarifying the function of the protein encoded by the cDNA. The full length clones were obtained by construction of full length enriched cDNA libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA easily without any special methods. The present sequence is a primer used to amplify a human cDNA clone provided in the invention
                                                                         Human; SPINK5; lympho-epithelial Kazal-type related inhibitor; LEKTI; se; serine protease inhibitor; atopic disease; Netherton's syndrome; asthma; eczema; hayfever; antiasthmatic; antiallergic; antiinflammatory; dermatological; PCR primer; sequencing primer; gene therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; full length cDNA; cDNA synthesis;
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                                                                                                                                                                                 SPINKS gene sequencing and PCR primer
                                                                                                                                                                                                                                                                       AAS44406;
                                                                                                                                                                                                                                                                                                               AAS44406 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 5 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 18; Page 132; 1380pp + Sequence Listing; English
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WO200164747-A1
                                     Homo sapiens.
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2000JP-00118774
2000JP-00183765.
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na T, Nagai
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Pred. No. 1.6e+03
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K, Kojima
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S, Otsuki
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RESULT 1311
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Best Local S
Matches 18
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15-MAR-2000;
23-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fragments of a SPINKS clone, sequencing primers and PCR primers for SPINKS. SPINKS encodes lympho-epithelial Kazal-type related inhibitor (LEKTI), a serine protease inhibitor. Susceptibility or predisposition to an atopic disease in a human subject can be detected by screening the genome for one or more polymorphic variants of SPINKS gene and/or expression of a variant LEKTI protein in a tissue. Carrier status of a subject or development of Netherton's syndrome is diagnosed by screening for the presence of loss-of-function mutations in the SPINKS gene. An expression vector comprising a nucleic acid encoding a serine protease inhibitor or its functional fragment can be used in screening for compounds with potential pharmacological activity by determining the serine protease activity of a protein previously identified as a ligand of the LEKTI protein. The atopic diseases include Netherton's Syndrome, asthma erzema and hauferner.
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03-MAR-2000;
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  Brooks-Wilson AR,
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2000GB-00005229.
                                                                         BRITISH COLUMBIA
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2000US-0213958P.
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No. 1.6e+03;
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     Clee
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disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a method for treating a patient diagnosed as having a lower than normal high density lipoprotein-cholesterol (HDL-C) level, a higher than normal triglyceride level, or a cardiovascular disease, involving administering a compound that modulates LXR- or RXR-mediated transcriptional activity or ABC1 expression or activity. The LXR gene product may be used in an assay to identify compounds useful for the treatment of a disease or condition selected a lower than normal HDL cholesterol level, a higher than normal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense oligonucleotide; groB; groEL; groES; inhibitor; growth; microorganism; Escherichia coli; Streptococcus pneumoniae; diagnosis; Streptococcus pyogenes; Staphylococcus aureus; Pseudomonas aeruginosa; antibacterial; antiviral; antiproliferative; antisense therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     S. aureus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Staphylococcus aureus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       microbial infection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         triglyceride level, and a cardiovascular disease
                                                                                                                                                                                                                                                                                                  WPI; 2001-355633/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-NOV-2000; 2000WO-CA001347.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200136625-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                    of microorganism, which hyts, useful to inhibit growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 659
                                                                                                                                                                                       antisense compounds targeting nucleic acid encoding groEL or of microorganism, which hybridize with and inhibit expression, useful to inhibit growth of microorganism having the genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                                                                      Į,
                                                                                                                                                                                                                                                                                                                                                                                                       GENESENSE TECHNOLOGIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 a lower than normal high density lipoprotein-cholesterol (HDL-C) higher than normal triglyceride level, or a cardiovascular by administering a compound that modulates LXR- or RXR-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GTGGCGCAATCTTGGCTCAC 678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   groE
                                                                                                                                                                                                                                                                                                                                                Young
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0166249P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   operon antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 A; 7 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 activity.
                                                                                                                                                                                                                                                                                                                                                      ¥
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.7%;
                                                                                                                                                                                                                                                                                                                                                   Dugourd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                      Ö
                                                                                                                                                                                                                                                                                                                                                                                                          INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16.8;
No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ۲.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO:425
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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                                                                                                                                                                                                                     groES
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The present invention specifically claims AAH56368 antisense oligonucleotides to nucleotide sequences generally, antisense compounds (I) comprising antis of 5-50 bases targeted to a nucleotide sequence enc

antisense

to AAH56832 which are encoding groE. More sense oligonucleotides groEL

encoding

English

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RESULT 1313
AAH56775
ID AAH5677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          therapeutics, prophylaxis and as research reagents and kits, e.g., to prevent or delay microbial infections in humans. They are also useful as molecular weight markers. AAH56362 to AAH56367 and AAH56833 to AAH56854 represent PCR primers for groß sequences which are used in the exemplification of the present invention. AAH56855 to AAH56870 represent groß nucleotide sequence given in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression of GL or GS, is claimed. (I) have antibacterial, antiviral and antiproliferative activities, and can be used in antisense therapy and for inhibiting expression of GL or GS in cells or tissues in vitro. (I) are also useful for inhibiting the growth of a microorganism, or inhibiting the expression of GL or GS gene in a microorganism (a bacterial cell or a virus) having a GL or GS gene which involves administering to the microorganism or to a cell infected with the microorganism, (I). (I) are also useful for treating a mammalian pathological condition mediated by the microorganisms having a pathological condition mediated by microorganisms having a GL or GS gene and administering (I) such that the growth of microorganism is inhibited. The antisense compounds are utilized for diagnostics,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       microorganism, where the antisense compound is complementary to GL or GS of a microorganism and specifically hybridises with and inhibits the expression of GL or GS, is claimed. (I) have antibacterial, antiviral and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense oligonucleotide; groE; groEI; groES; inhibitor; growth; microorganism; Escherichia coli; Streptococcus pneumoniae; diagnosis; Streptococcus pyogenes; Staphylococcus aureus; Pseudomonas aeruginosa; antibacterial; antiviral; antiproliferative; antisense therapy;
                                                                                                                                                                             18-NOV-1999;
                                                                                                                                                                                                                                      20-NOV-2000; 2000WO-CA001347
                                                                                                                                                                                                                                                                                                      25-MAY-2001
                                                                                                                                                                                                                                                                                                                                                           WO200136625-A2
                                                                                                                                                                                                                                                                                                                                                                                                                       Staphylococcus aureus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       microbial infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     S. aureus groE operon antisense oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH56775
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH56775 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   shock protein (HSP)60) (GL) and groES (HSP10) (GS) gene from
                                                     Wright JA,
WPI; 2001-355633/37.
                                                                                                                     (GENE-) GENESENSE TECHNOLOGIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          601
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TTTTTATTTTTAATTTTTTG 620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TTTTTATTTCAACTTTTTG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                     Young AH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                             99US-0166249P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 C; 1 G; 14 T; 0 U; 0 Other;
                                                     Dugourd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 16.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NO:423
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Novel antisense compounds targeting nucleic acid encoding groEL or gene of microorganism, which hybridize with and inhibit expression genes, useful to inhibit growth of microorganism having the genes.

of the

Novel antisense compound 8-30 nucleobases acid molecule encoding human mdm-2 useful

The present invention specifically claims AAH56368 to AAH56832 which

are

in length targeted

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33; 77pp; and reducing

English

hyperproliferation

in length targeted to a nucleic for modulating the expression ation of human cells.

The present invention relates to an antisense compound 8-30 nucleobases in length targeted to nucleobases 1-308 of the 5' untranslated region,

Page 52; 110pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cc microorganism, where the antisense compound is complementary to GL or GS of a microorganism and specifically hybridises with and inhibits the CC expression of GL or GS, is claimed. (I) have antibacterial, antiviral and CC antiproliferative activities, and can be used in antisense therapy and CC inhibition of expression of GL or GS in cells or tissues in vitro. (I) are CC inhibiting expression of GL or GS in cells or tissues in vitro. (I) are CC also useful for inhibiting the growth of a microorganism, or inhibiting CC the expression of GL or GS gene in a microorganism, or inhibiting CC the expression of GL or GS gene which involves administerial cell or a CC virus) having a GL or GS gene which involves administering to the CC microorganisms or to a cell infected with the microorganism, (I). (I) are CC also useful for treating a mammalian pathological condition mediated by the microorganisms which involves identifying a eukaryotic organism CC daving a pathological condition mediated by microorganisms having a GL or CC gene and administering (I) such that the growth of microorganism is CC inhibited. The antisense compounds are utilised for diagnostics, to prevent or delay microbial infections in humans. They are also useful as CC molecular weight markers. AAH5632 to AAH56357 and AAH56831 to AAH56854 CC exemplification of the present invention. AAH56855 to AAH56870 represent CC groß mucleotide sequence given in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1314
AAF80889/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antisense oligonucleotides to nucleotide sequences encoding groß. More generally, antisense compounds (I) comprising antisense oligonucleotides of 5-50 bases targeted to a nucleotide sequence encoding großL (heat shock protein (HSP)60) (GL) and großS (HSP)60) (GS) gene from a shock protein (HSP)60) (GL) and großS (HSP)60) (GS) gene from a shock protein (HSP)60) (GL) and großS (HSP)60) (GS) gene from a shock protein (HSP)60) (GL) and großS (HSP)60) (GS) gene from a shock protein (HSP)60) (GL) and großS (HSP)60) (GS) gene from a shock protein (HSP)60) (GL) and großS (HSP)60) (GS) gene from a shock protein (HSP)60) (GL) and großS (HSP)60) (GS) gene from a shock protein (HSP)60) (GL) and großS (HSP)60) (GS) gene from a shock protein (HSP)60) (GS) and großS (HSP)60) (GS) gene from a shock protein (HSP)60) (GS) and großS (HSP)60) (GS) gene from a shock protein (HSP)60) (GS) and großS (HSP)60) (GS) gene from a shock protein (HSP)60) (GS) and großS (HSP)60) (GS) gene from a shock protein (HSP)60) (GS) and großS (HSP)60) (GS) gene from a shock protein (HSP)60) (GS) gene from a shock protein (HSP)60) (GS) gene from a shock protein (HSP)60) (HSP)60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense; mdm2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human mdm2 phosphorothioate oligonucleotide #263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF80889
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF80889 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                          WPI; 2001-190948/19.
                                                                                                                                                                                                                                                                                                                                Miraglia LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-FEB-2001.
                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              604 TTATTTTAATTTTTGAGA 623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTATTTCAACTTTTGAGA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                               PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                Nero P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-00048810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-00280805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hyperproliferation; cancer; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 C; 2 G; 11 T; 0 U;
                                                                                                                                                                                                                                                                                                                                      Graham MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 16.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                      Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 Other;
                                                                                                                                                                                                                                                                                                                                      Cowsert LM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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RESULT 1315
AAF80875/C
ID AAF8087
XX AAF8087
XX DE Human T
XX ANTISEN
XX ANTISEN
XX HOMO S8
XX HOMO S8
XX HOMO S8
XX O6-FEB-
XX O6-FEB-
XX (ISIS-)
XX MOVEl:
PT acid mc
PT acid 
RESULT 1316
AAF80882/c
ID AAF8088
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Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to an antisense compound 8-30 nucleobases in length cargeted to nucleobases 1-308 of the 5' untranslated region, 1776-1806 of the translation termination codon region or 1818-2370 of the 3' untranslated region of a nucleic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro. The hyperproliferative disorder includes cancer or psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel antisense compound 8-30 nucleobases in length targeted to a nucleobased molecule encoding human mdm-2 useful for modulating the expression of human mdm-2 and reducing hyperproliferation of human cells.
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  AAF80882 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                    Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-FEB-2001.
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                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                               TCAGCCTCCCAAGTAGCTGG 562
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                                                                                                                                                                                                                                                                                                                                    BP; 6 A; 3 C; 7 G; 4 T; 0 U; 0 Other;
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Pred.
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Pred. No. 1.6e+03
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No. 1.6e+03
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RESULT 1317
AAF80886/c
ID AAF8088
XX AAF8088
XX AAF8088
XY CAAF8088
XY CAAF808
XY CAA
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Best Local Similarity
Matches 18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF80886 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of human mdm-2 and reducing hyperproliferation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-190948/19.
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                                                                                                                                                                                                                                                                                                                                                       Human mdm2
                                                                                                                                                                                                                                                                                                                                                                                                           02-MAY-2001
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                                                    26-MAR-1998;
                                                                                                     26-MAR-1999;
                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                           Antisense;
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                                                                                                                                                                                                                                                                                                        hyperproliferation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                           cancer; psoriasis;
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(ISIS-) ISIS

PHARM INC

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RESULT 1318
AAS09243
ID AAS0924
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AC AAS0924
XX
AC AAS0924
XX
DT 24-OCT-
DE PCR pCR pCR
XX
Human;
KW Human;
KW PCR p7
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POS Homo (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-MAY-1992;
16-APR-1993;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1776-1806 of the translation termination codon region or 1818-2370 of the 3' untranslated region, of the attranslated region of a nucleic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro. The hyperproliferative disorder includes cannot be represented by the superproliferative disorder includes cannot be required to the superproliferative disorder to the superproliferative disorder to the superproliferative disorder to the superproliferative disorder to the super
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel antisense compound 8-30 nucleobases in length targeted to a nucleic acid molecule encoding human mdm-2 useful for modulating the expression of human mdm-2 and reducing hyperproliferation of human cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Miraglia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; familial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer #1 for marker D9S58 associated with familial dysautonomia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAS09243 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-190948/19
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                                                                                                                                                                                                                                                                                                                                                                                      polymorphisms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Blumenfeld
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US6262250-B1
                                                                                                                                                                                                                                                Disclosure;
                                                    The present sequence for PCR primer #1 is used with PCR p (AAS09244) to amplify DNA marker D9S58. Various oligonucl (AAS09239-AAS09272) are described in an invention relating the control of the control
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GEHO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                 for detecting presence of polymorphisms linked to gene associated h familial dysautonomia (FD), comprises specific primers which detect ymorphisms, D9S309 and D9S310 identified in candidate region for FD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                    601
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93US-00049678.
95US-00480655.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-00455683
                                                                                                                                                                                                                                                10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          dysautonomia; chromosome 9q31-q33; Riley-Day syndrome;
al loss of neuron; nervous system; DNA marker D9S58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'n,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Graham MJ,
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                                                                                                                                                                                                                                                English.
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Pred. No. 1
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              familial dysautonomia (FD)
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                                                  markers in 26 FD families. A kit to detect the presence of polymorphisms linked to a gene associated with FD, the Riley-Day syndrome (an autosomal recessive disorder characterised by developmental loss of neurons from sensory and autonomic nervous system) in an individual, comprises a nucleic acid primer of at least 15 contiguous nucleotides and at least one other reagent. The kits are useful for diagnosing familial dysautonomia and the test can be used prenatally to screen a foetus, or presymptomatically to screen a subject at risk in affected FD families
Sequence 20
                                                                                                                                                                                                                                                                                                The FD gene has been mapped to chromosome 9q31-q33 by linkage with 10 DNA
   BP; 5 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
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Query Match
Best Local S
Matches 18
                725 CCTGAGTAGCTGGGACTACA 744
                                   18;
                                           Similarity
CCTGAGTAGCCGGGACTATA 20
                                    Conservative
                                           1.7%;
                                   <u>,</u>
                                            Pred.
                                                      Score
                                    Mismatches
                                             16.8;
No. 1
                                             .6e+03
                                                      DB 1;
                                                      Length
                                     Indels
                                    0
                                     Gaps
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SNP specific upper PCR primer SEQ ID 337.
                     14-AUG-2001
                                         AAH37541;
                                                              AAH37541
                                                              standard;
                      (first
                                                              DNA;
                      entry)
                                                              20
                                                               ВÞ
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Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.

Homo sapiens.

WO200129262-A2

26-APR-2001.

13-OCT-2000; 2000WO-US028436

15-OCT-1999; 99US-0160096P

(ORCH-) ORCHID BIOSCIENCES INC.

Picoult-Newburg ۲ Poh1 Σ,

WPI; 2001-290930/30

acid New genotyping oligonucleotide, absence or identity of single po e, useful for detecting the polynucleotide polymorphism m in a nuc

Claim 1; Page 51; 83pp; English.

primer extension (SNPB) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide

of polymorphisms associated with

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RESULT 1320
AAH24276/c
ID AAH2427
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Best Local S
Matches 18
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comprising a recombinant expression construct encoding a mammalian p21 on p16 gene. The p21 and p16 proteins are cyclin dependent kinase (CDK) inhibitors cause cell cycle arrest in a variety of physiological situations, p21 and p16 are intimately associated with the process of sensence in mammalian cells. The invention also encompasses cells additionally containing a construct comprising a reporter gene under the control of a promoter from a mammalian gene whose expression is induced or inhibited by a CDK inhibitor such as p21 or p16. The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200138532-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH24276 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 7 A; 7 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                   New recombinant mammalian fibrosarcoma cell useful for identifying compounds that inhibit CDK inhibitor-mediated modulation of cellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-NOV-1999; 99US-00449589.
07-APR-2000; 2000WO-US009286.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cell cycle arrest; drug screening; senescence marker;
senescence inhibitor; potentiator; gene expression modulator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human PLK1 promoter; p21-repressible promoter; reporter construct; recombinant fibrosarcoma cell; p21 expression construct; p16; CDK inhibitor; cyclin dependent kinase inhibitor; cellular senescence;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH24276;
                                                                                                                                                                                                                                      The invention relates to a recombinant mammalian fibrosarcoma
                                                                                                                                                                                                                                                                                               Example 6; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-OCT-2000; 2000WO-US028082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cellular proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ъ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GATGGGGTTTCACCATGTTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Roninson IB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oliferative disorders; cancer; age-related disease; disease; amyloidosis; atherosclerosis; arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ILLINOIS FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first
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                                                                                                                                                                                                                                                                                            72; 136pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score
Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
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     The invention
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Query Match Best Local Similarity

1.7**%**; 90.0**%**; 5 C; 7 G;

Pred. No.

1.6e+03;

16.8;

DB 1;

Length

Sequence

20

B₽;

5 A;

3 T; 0 U; 0 Other;

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RESULT 1321
AAF29798
ID AAF2979
XX AAF2979
XX O9-APR-
XX Preseni
XX Human;
KW Preseni
XX Homo si
PN WO2000
XX Homo si
YX WO21010
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YX WO2000
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        additionally relates to the identification of genes whose expression is modulated by a CDK inhibitor. Such genes (which include connective tissue growth factor, serum amyloid A, integrin beta-3, activin A, natural killer cell protein 4, Mac2 binding protein and tissue transglutaminase) can be used as markers of cellular senescence. Recombinant cells of the invention are used to identify compounds which inhibit, promote or potentiate senescence, or which modulate the effects of CDK inhibitor-mediated induction or repression of gene expression. Compounds identified using methods of the invention may be used in the treatment of cellular proliferative disorders such as cancers, or age-related diseases such as Alzheiner's disease, amyloidosis, atheroselerosis, and arthritis. Sequences AAH24276- AAH24277 represent PCR primers used in an exemplification of the invention to amplify the p21-repressible human pLKI promoter for construction of a p21-responsive reporter construct
                                                             The present invention describes a method for determining the presence or susceptibility to Alzheimer's disease in humans, involving detecting genetic lesion in the presentiner—1 (PSENI) gene, found on chromosome The genetic lesion is a polymorphism in the promoter or upstream regulatory region of the gene. The invention also describes transgenic animals which can be used to identify compounds useful in treating
                                                                                                                                                                                                                                        Determining wether a human subject has or is at -onset) Alzheimer's disease comprises detecting genetic lesion in the presention-1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200079000-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              preseniline-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; PSEN1; Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Preseniline-1
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                                                                                                                                                                                                                                                                                                                                                                Theuns
                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                         (VLAA-) VLAAMS INTERUNIVERSITAIR INST BIOTECHNOG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCAAATGCTGGGATTACAGG 1
                                                                                                                                                                                                        Page 45; 56pp; English.
                                               disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chromosome
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          promoter PCR primer Prom23F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
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                                                                                                                                                                                                                                                                                                                                                                  Van
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                                                                                                                                                                                                                                                                                                                                                                    Broeckhoven
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease; polymorphism; diagnosis;
14; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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Pred. No. 1.
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                                                                 so describes transgenic useful in treating
                                                                                                                                                                                                                                                                 risk of developing (early the presence/absence of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
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                                                                                                                             detecting a
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RESULT 1322
AAC67119/c
ID AAC6711
RESULT 1323
AAS21753/c
ID AAS21753 &
XX AAS21753;
AC AAS21753;
XX DT 21-NOV-200
XX Mouse Surv
XX
                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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07-OCT-1994;
08-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC67119 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Determining predisposition of a human to hypertension, involves analyzing DNA sequence of angiotensinogen for a mutation which is in linkage disequilibrium with specific mutation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-APR-2001
                                                                                                                                                                                                                                                                                                                                               The present invention describes a method for determining the predisposition of an individual to hypertension, involving analysing angiotensinogen (AGT) alleles they possess. Individuals with a M235T mutation in the angiotensinogen gene are at an increased risk of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-101691/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jeunemaitre
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       predisposition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Angiotensinogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human growth
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                 Mouse Survivin antisense oligonucleotide
                                                                                                                                                                                                                                                                                                      Sequence
                                                21-NOV-2001
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INSERM INST NAT SANTE &
                                                                                                                                                                                                                                                                                                                                                                                                                            3; Col 12; 26pp; English.
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                                                                                                          standard;
                                                                                                                                                                                                CACCCAGGCTGGAGTGCAGT 660
                                                                                                                                                                                                                                                                                                      20 BP; 4 A; 8 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                  CTCCGAGGCTGGAGTGCAGT 1
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                                                                                                                                                                                                                                              Conservative
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                                                (first entry)
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94US-00319545.
98US-00092988.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AGT; variant; human; hypertension; M235T mutation;
                                                                                                          DNA;
                                                                                                                                                                                                                                                          1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene
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                                                                                                            ВP
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                                                                                                                                                                                                                                                          Score 16.8;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kotelevtsev
                                                                                                                                                                                                                                                          1.6e+03
                                                                                                                                                                                                                                                                           DB 1;
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AAS29504/c
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         antisense oligonucleotides are used in the treatment of an animal suffering from a disease or condition associated with Survivin, e.g. a CC hyperproliferative condition such as cancer, and comprises administering CC a therapeutically or prophylactically effective amount of the antisense CC oligonucleotide so that expression of Survivin is inhibited. The CC oligonucleotide so a laso be used to treat a human suffering from a CC disease or condition characterised by a reduction in apoptosis comprising CC administering the antisense oligonucleotide to a human. In addition, the antisense oligonucleotide to a human. In addition, the care and a cytotoxic chemotherapeutic agent e.g. taxol or cisplatin, can be used to modulate apoptosis, cytokinesis or the CC cell cycle, or inhibit the proliferation in a cancer cell by contacting the cell with the antisense oligonucleotide. AAS21521-AAS21768 represent Survivin nucleic acids, and antisense oligonucleotides targeted to Survivin, used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        acid molecule encoding human Survivin, where the antisense oligonucleotide inhibits the expression of human Survivin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Survivin; human; mouse; cytostatic; antisense oligonucleotide; hyperproliferative condition; cancer; apoptosis; cytokinesis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   treatment of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-FEB-2000; 2000US-00496694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JAN-2001; 2001WO-US002939
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 18; Page 62; 120pp;
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                    modified_base
                                                                                              Human; mdm2; hyperproliferative disorder; cancer; psoriasis; atherosclerosis; tumour; cytostatic; anti psoriatic; anti arteriosclerotic; vasotropic; antisense; phosphorothioa
                                                                                                                                                        Human mdm2 antisense oligonucleotide
                                                                                                                                                                                                                                              AAS29504 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
                                                                   Homo sapiens
                                                                                                                                                                                       21-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense compounds for modulating
                                                                                                                                                                                                                                                                                                                          20 TANAGGTGTGAGCCACCACG
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                                                                                                                                                                                       (first entry)
      Location/Qualifiers
1. .20
/*tag= a
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KW Human;
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                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to antisense compounds, 8-30 nucleobases in CC length targeted to the 5' untranslated region, translation termination CC codon region, 3' untranslated region, coding region or translation start CC site of a nucleic acid encoding human mdm2, where the antisense compound modulates the expression of human mdm2. The antisense oligonucleotides of CC the invention are useful for encoding human mdm2 and for inhibiting the CC expression of human mdm2. They may be used for treating an animal having CC a disease or condition associated with amplification of mdm2 gene or overexpression of mdm2 e.g. a hyperproliferative disorder such as cancer (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis, CC fibrosis, atheroselerosis or restenosis, tumours, colorectal carcinoma CC and chronic myelogenous leukemia. The antisense compound may be CC administered with a chemotherapeutic agent to overcome drug resistance. CC The antisense compound reduces hyperproliferation of human cells. The method, which involves the use of the antisense compound, is also useful CC for detecting the role of mdm2 expression in various cell functions and CC diagnostic tools. AAS29242-AAS29507 represent the human mdm2 antisense CC oligonucleotides of the present invention
                                                                                                                                                                                     RESULT 1325
                                                                                                                                                                                                                                                                                                         Matches 18;
                                                                                                                                                                                                                                                                                                                        Query Match
Best Local :
Human; mdm2; hyperproliferative disorder; cancer; psoriasis;
atherosclerosis; tumour; cytostatic; anti psoriatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             An antisense compound, useful for trea nucleobases targeted a region (e.g. tr of a nucleic acid encoding human mdm2.
                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 2 A; 3 C; 11 G; 4 T; 0 U; 0 Other;
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26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2001016575-A1.
                                                Human mdm2 antisense oligonucleotide 31469.
                                                                                                                                                      AAS29490 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 9; Page 18; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-JAN-2001; 2001US-00752983
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                                                                                   21-NOV-2001
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GRAHAM M J.
MONIA B P.
                                                                                                                                                                                                                                                                                                                        Similarity
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COWSERT L M.
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                                                                                                                                                                                                                                      GCCCACCTCGGCCTCCCAAA 1
                                                                                                                                                                                                                                                                                                         Conservative
                                                                                   (first entry)
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99US-00280805
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additionally bases 1-6 and bases 15-20 are 2'-0-
methoxyethyl bases, and bases 7-14 are deoxynucleotides"
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                                                                                                                                                      20 BP
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0; Mis
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 for treating e.g. cancer, comprises (e.g. translation termination codon
                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                      Length 20;
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RESULT 1326 AAS29497/c ID AAS2949 XX AC AAS2949

AAS29497 standard;

DNA;

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AAS29497;

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The present invention relates to antisense compounds, 8-30 nucleobases in clearly thargeted to the 5' untranslated region, translation termination codon region, 3' untranslated region, coding region or translation start site of a nucleic acid encoding human mdm2, where the antisense compound modulates the expression of human mdm2. The antisense oligonucleotides of the invention are useful for encoding human mdm2 and for inhibiting the expression of human mdm2. They may be used for treating an animal having a disease or condition associated with amplification of mdm2 gene or overexpression of mdm2 e.g. a hyperproliferative disorder such as cancer (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis, fibrosis, atherosclerosis or restenosis, tumours, colorectal carcinoma and chronic myelogenous leukemia. The antisense compound may be administered with a chemotherapeutic agent to overcome drug registance. The antisense compound reduces hyperproliferation of human cells. The method, which involves the use of the antisense compound, is also useful physiological processes and useful in both clinical research and antisense coll functions and diagnostic tools. AAS2944-AAS29507 represent the human mdm2 antisense
                                                                                                                          Query Match
Best Local Similarity
                                                                                                    Matches
                                                                                                                                                                                                      Sequence 20 BP; 6 A; 3 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 18; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of a nucleic acid encoding human
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(GRAH/)
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26-MAR-1999;
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                                                                                                                                                                                                                                                          oligonucleotides of the present invention
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COWSERT L M.
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GRAHAM M J.
                                      TCAGCCTCCCAAGTAGCTGG 562
                                                                                                    Conservative
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99US-00280805.
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additionally bases 1-6 and bases 15-20 are 2'-0-
methoxyethyl bases, and bases 7-14 are deoxynucleotides"
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                                                                                                                       1.7%;
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Pred. No. 1.
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                                                                                                       Mismatches
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translation
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termination codon region)
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                                                                                                       Indels
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The present invention relates to antisense compounds, 8-30 nucleobases in CC length targeted to the 5' untranslated region, translation termination CC codon region, 3' untranslated region, coding region or translation start CC site of a nucleic acid encoding human mdm2, where the antisense compound CC modulates the expression of human mdm2. The antisense oligonuclectides of CC the invention are useful for encoding human mdm2 and for inhibiting the CC expression of human mdm2. They may be used for treating an animal having CC disease or condition associated with amplification of mdm2 gene or CC (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis, CC fibrosis, atherosclerosis or restenosis, tumours, colorectal carcinoma CC and chronic myelogenous leukemia. The antisense compound may be CC administered with a chemotherapeutic agent to overcome drug resistance. The antisense compound reduces hyperproliferation of human cells. The CC method, which involves the use of the antisense compound, is also useful CC for detecting the role of mdm2 expression in various cell functions and CC diagnostic tools. AAS29242-AAS29507 represent the human mdm2 antisense CC cligonucleotides of the present invention
   á
                                  Query Match
Best Local S
Matches 18
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atherosclerosis; tumour; cytostatic; anti psoriatic;
anti arteriosclerotic; vasotropic; antisense; phosphorothioate; ss
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26-MAR-1999;
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                                                                                               Sequence
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      316
                                    l Similarity
18; Conser
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COWSERT L M.
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GTAGAAACAGGGTTTCACTG 335
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                                    Conservative
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99US-00280805.
                                                                                                 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/mod="0THER= All phosphorothioate linkages,
note="0THER= All phosphorothioate linkages,
additionally bases 1-6 and bases 15-20 are 2'-0-
methoxyethyl bases, and bases 7-14 are deoxynucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cocation/Qualifiers
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                                                                                                  7
                                                                                                 C; 4
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                                                                                                 G;
                                                    Score 16.8;
Pred. No. 1
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                                       Mismatches
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                                                      .6e+03
                                                                                                  0 Other;
                                                                     DB 1;
                                                                 Length 20;
                                       Indels
                                     0;
                                       Gaps
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20

GTAGAGACAGGGTTTCACCG 1

20

BP; 6 A;

4 C;

8 G;

N Ţ;

0 U; 0 Other

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The present invention relates to antisense compounds, 8-30 nucleobases in CC length targeted to the 5' untranslated region, translation termination CC codon region, 3' untranslated region, coding region or translation start CC site of a nucleic acid encoding human mdm2, where the antisense compound CC modulates the expression of human mdm2. The antisense oligonucleotides of CC the invention are useful for encoding human mdm2 and for inhibiting the CC expression of human mdm2. They may be used for treating an animal having CC adisease or condition associated with amplification of mdm2 gene or CC (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis, cC fibrosis, atheroselerosis or restenosis, tumours, colorectal carcinoma CC and chronic myelogenous leukemia. The antisense compound may be administered with a chemotherapeutic agent to overcome drug resistance. CC The antisense compound reduces hyperproliferation of human cells. The CC method, which involves the use of the antisense compound, is also useful CC for detecting the role of mdm2 expression in various cell functions and CC diagnostic tools. AAS29242-AAS29507 represent the human mdm2 antisense CC oligonucleotides of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; mdm2; hyperproliferative disorder; cancer; psoriasis; atherosclerosis; tumour; cytostatic; anti psoriatic; anti arteriosclerotic; vasotropic; antisense; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human mdm2 antisense oligonucleotide 31629.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAS29501;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAS29501 standard;
                                                                                                                                                                                                                                                                                                                                                                                 An antisense compound, useful for treating e.g. nucleobases targeted a region (e.g. translation of a nucleic acid encoding human mdm2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Miraglia LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MIRA/) MIRAGLIA L J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-JAN-2001; 2001US-00752983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GRAH/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ) GRAHAM M J.
) MONIA B P.
) COWSERT L M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NERO
                                                                                                                                                                                                                                                                                                                                                    Page 18; 81pp;
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99US-00280805
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                     English.
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1-6 and bases 15-20 are 2'-0-
, and bases 7-14 are deoxynucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Monia
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                                                                                                                                                                                                                                                                                                                                                                                                     cancer, comprises
termination codon region)
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RESULT 1328
ABZ72236
ID ABZ7223
XX Human;
XX WO20017
XX WO20017
XX I3-APR-
XX (GENO-)
XX I3-APR-
XX (GENO-)
XX WPI; 20
DR WPI; 20
DR
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                                                                                                                                                                                                                                  The invention relates to isolated genes (Gene 216) from human chromosome CC 20p13-p12 and the proteins they encode. The nucleic acids and proteins CC may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate Gene 216 expression. For example, the conclusic acids (or vectors) and protein may be used to treat disorders CC associated with decreased expression by rectifying mutations or deletions CC in a patient's genome that affect the activity of gene 216 by expressing CC inactive proteins or to supplement the patients own production of Gene C16 proteins. Additionally, the nucleic acids may be used to produce the secreted Gene 216 protein, by inserting the nucleic acids into a host C complementary sequences may also be used as DNA probes in diagnostic CC assays to detect and quantitate the presence of similar nucleic acids and CC complementary sequences may also be used as DNA probes in diagnostic CC assays to detect and quantitate the presence of similar nucleic acid complements of Gene 216 ence 116 protein may also be used as antigens in CC the production of antibodies against Gene 216 and in assays to identify CC modulators of Gene 216 expression and activity. The anti-Gene 216 antibodies and antigens that may be in samples (e.g. apents for detecting the presence of Gene 216 proteins in samples (e.g. by enzyme linked immunosorbant assay or ELISA). Disorders that may be consented. diagnosetic and antigens be the activity of gene 210 and in absorbant may be consented.
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated genes (Gene 216) from human chromosome 20p13-p12 and the proteins they encode, useful for the prevention, diagnosis and treatment of asthma, obesity and inflammatory house diagnosis.
   prevented, diagnosed and/or treated by the above methods include, for example asthma, obesity and inflammatory bowel disease. The present sequence is that of a Gene 216 related primer used in examples of the invention. The primers are used in the physical mapping of the gene (ABZ72067-ABZ72088), polymorphism identification using single strand conformational polymorphism (SSCP) analysis (ABZ7091-ABZ72184), sequencing (ABZ72185-ABZ72268) and genotyping (ABZ72317-ABZ72362)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; Gene 216; chromosome 20p13-p12; antiasthmatic; anorectic; antiinflammatory; gastrointestinal; gene therapy; vaccine; asthm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gene 216 SSCP
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Pred. No. 1.
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The invention relates to antisense compounds targetted to nucleic acid encoding RECQL2 (gene associated with Bloom's disorder) to inhibit the expression of RECQL2. Antisense compounds of the invention are useful f treating diseases associated with expression of RECQL2, in humans. They are useful for diagnostics, therapeutics and as research reagent, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 4 A; 7 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                Antisense compounds targeted to nucleic acids encoding RECQL2 associated with Bloom's disorder, for modulating RECQL2 expression and treating diseases e.g. tumors associated with expression of the RECQL2 in humans.
                                                                                                                                                                                                  WPI; 2002-535979/57
                                                                                                                                                                                                                                                                                                                                01-MAR-2001; 2001US-00798096
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense; RECQL2; Bloom's disorder; prophylaxis; infection; tumour; inflammation; therapy; human; phosphorothioate; ss.
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                                                                                                  44; 86pp;
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formation. They are also useful in antisense therapy.
sequence is an antisense oligonucleotide targetted to
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The present invention relates to antisense oligonucleotides and methods for modulating the expression of human or mouse Protein Phosphatase 2 catalytic subunit alpha. The antisense oligonucleotides are useful for inhibiting the expression of Protein Phosphatase 2 catalytic subunit alpha and for treating diseases or conditions associated with aberrant expression of Protein Phosphatase 2 catalytic subunit alpha. Such diseases include diabetes and cancer. The antisense oligonucleotides are also useful for diagnostics, therapeutics, and prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. They are also useful as research reagents for distinguishing between functions of various members of a biological pathway. ABS5400-ABS65477 represent human or mouse Protein Phosphatase 2 catalytic subunit alpha antisense oligonucleotides which comprise a phosphorothioate backbone
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Pred. No. 1
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                                 The present invention relates to antisense oligonucleotides and methods for modulating the expression of human or mouse casein kinase 2-alpha prime. The antisense oligonucleotides are useful for inhibiting the expression of casein kinase 2-alpha prime, and for treating diseases or conditions associated with aberrant expression of casein kinase 2-alpha prime. Such diseases include diabetes mellitus, and hyperproliferative disorders (particularly cancers e.g. breast cancer, prostate cancer, or liver cancer). The antisense compounds are also useful for diagnostics, therapeutics, prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits, and in distinguishing between functions of various members of a biological pathway. ABS67840-ABS67917 represent human or mouse casein kinase 2-alpha prime antisense oligonucleotides which comprise a phosphorothicate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hyperproliferative disorder; breast cancer; prostate cancer; liver cancer; infection; inflammation; tumour formation; cytostatic; antidiabetic; antiinflammatory; antimicrobial; phosphorothioate;
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                                                                                                                                                                                                                                                Claim 3; Page 94; 129pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diabetes mellitus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .6e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2;
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AGTGGCGCAATCTCAGCTCA 1

15-NOV-2002 (first entry)

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RESULT 1332
ABS67843/C
ID ABS6784
AC ABS6784
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X S X
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                                                                                                            RESULT 1333
ABS52459
                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to antisense oligonucleotides and methods for modulating the expression of human or mouse casein kinase 2-alpha prime. The antisense oligonucleotides are useful for inhibiting the expression of casein kinase 2-alpha prime, and for treating diseases or conditions associated with aberrant expression of casein kinase 2-alpha prime. Such diseases include diabetes mellitus, and hyperproliferative disorders (particularly cancers e.g. breast cancer, prostate cancer, or liver cancer). The antisense compounds are also useful for diagnostics, therapeutics, prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits, and in distinguishing between functions of various members of a biological pathway. ABS67840-ABS67917 represent human or mouse casein kinase 2-alpha prime antisense oligonucleotides which comprise a phosphorothioate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; casein kinase 2-alpha prime; diabetes mellitus; hyperproliferative disorder; breast cancer; prostate cancer; liver cancer; inflammation; tumour formation; cytostatic; antidiabetic; antiinflammatory; antimicrobial; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABS67843 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-FEB-2002; 2002WO-US002772.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human casein kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABS67843
                        ABS52459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense oligonucleotides targeted to nucleic acid encoding casein kinase 2-alpha prime, useful for diagnosing and/or treating a disease or condition associated with expression of casein kinase 2-alpha prime.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-FEB-2001; 2001US-00780173
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                                                                                   ABS52459 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-627539/67.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                               993
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                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page
                                                                                                                                                                                                                                                                                            CCCGGGCTCAAGCGATTCTC 1012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B₽;
                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94; 129pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Þ
                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2-alpha prime antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                  1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wyatt JR;
                                                                                   20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7
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                                                                                   ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G; 3 T;
                                                                                                                                                                                                                                                                                                                                                                                  Score 16.8; DB 1;
Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 20;
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CC fidelity to the Li consensus sequence in their 5 or 3 regulatory region (Li= long interspersed nuclear element, LIME-1). Also included are (1) cc identifying an individual at risk for or suffering from a complex disease (CC comprises: (a) identifying or detecting the amount of intronic regions of genes containing full length Li elements or in 5 or 3 regulatory (CC regions of genes containing a full length high fidelity consensus Li (CC sequence of the individual's DNA from a sample; and (b) comparing the (CC genes or the 5 or 3 regulatory regions with a control sample of DNA (CC genes or the 5 or 3 regulatory regions with a control sample of DNA (CC suffering from a complex disease, where the genes identified are involved (CC suffering from a gent selected from an Li antisense oligonucleotide, an antibody directed against Li mRNA, and an antibody directed against Li mRNA, and an antibody directed against Li mRNA, and an antibodies or auto (CC antibodies directed against ribonucleo-protein particles having Li mRNA (CC complements, Li DNA, mRNA or protein products which indicates that the control sate of the complex disease, where the genes is a trisk for or suffering from a complex disease (CC antibodies directed against ribonucleo-protein particles having Li mRNA (CC complements, Li DNA, mRNA or protein products which indicates that the control complex disease, autoimmune disease, soft complex disease (CC altheimer's disease, autoimmune diseases, schazophrenia, systemic lupus (CC artibodies or multiple coloronic disease).
                                                                           erythematosus, multiple sclerosis, insulin-dependent diabetés mellitus, rheumatoid arthritis, phemphigus, psoriasis, autoimmune thyroid disease, scleroderma, mixed connective tissue disease, polymyositis, dermatomyositis, Sjogren's syndrome, pemphigoid, vitiligo, primary biliary cirrhosis, chronic active hepatitis, Crohn's disease, ulcerative colitis and pernicious anaemia). Detection of the protein products of L1 elements, either ORF1/P40 or ORF2 gene products can be used to indicate the presence in cells, tissue, or body fluids of potential immune system triggers that can induce or exacerbate autoimmune disease. The present sequence is a PCR primer included in the sequence listing but not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identifying a gene involved in a complex disease, e.g. schizophrenia, comprises detecting genes having full-length L1 element in their intronic region or high sequence fidelity to L1 consensus sequence in the 5' or 3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   psoriasis; autoimmune thyroid disease; polymyositis; vitiligo; mixed connective tissue disease; dermatomyositis; Sjogren's synd pemphigoid; primary biliary cirrhosis; chronic active hepatitis; Crohn's disease; ulcerative colitis; pernicious anaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ss; long interspersed nuclear element; LINE-1; p40; PCR; primer; ORF2; L1; Alzheimer's disease; autoimmune disease; schizophrenia; systemic lupus erythematosus; multiple sclerosis; scleroderma; insulin-dependent diabetes mellitus; rheumatoid arthritis; phemph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-DEC-2001; 2001WO-US049353.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to identifying a gene involved in a complex disease comprising identifying genes containing full-length L1 elements in their intronic region or containing a full length L1 element with high sequence intronic region or containing a full length L1 element with high sequence fidelity to the L1 consensus sequence in their 5' or 3' regulatory region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 137; 138pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-643381/69
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-DEC-2000; 2000US-0256673P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       regulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (HOSP-) HOSPITAL FOR SPECIAL SURGERY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       region
                                                  anywhere
                                                    else
                                                    in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer
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Sequence

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                                                                                                                                                    CC The present invention describes a method of arraying genome clones. The CC multiwell plates (a) clones of the genomic libraries contained in CC multiwell plates; (b) a primer designed based on the chromosome marker CC sequence is added to the mixture to carry out an amplification reaction; (C) a signal corresponding to the marker is detected from the resultant CC amplified product to specify the discrimination Nos. of the multiwell CC plates containing the clones having said marker sequence; (d) the order CC of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to the multiwell CC plates; (e) the clones in the multiwell plates of the specified CC discrimination Nos. are mixed respectively in each wells of longitudinal CC discrimination Nos. are mixed respectively in each wells of longitudinal CC resultant cultures are amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell CC plates are specified from the detected result; and (i) the clones are enconstituted as the positions on the chromosome and arrayed. The CC primers for human chromosome 1936-35 DNA, and ABI45322 represent CC expresent PCR primers for human chromosome 21q22.1, which are CC specifically claimed for use in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 18; Conserv
                                                              Matches
                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arraying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-MAR-2000; 2000JP-00066716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-MAR-2001; 2001JP-00068285
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human chromosome 21q22.1 PCR primer SEQ ID NO:2440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABL45396;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABL45396 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 6; Page 53; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (RIKA ) RIKAGAKU KENKYUSHO. (GENO-) GENOTEX YG.
                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              199 ATGTTGGTCAGGCTGGTCTC 218
                              732
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                                                              18;
 _
                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genome clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATGTTGGCCAGGCTGATCTC 20
                                                                                                                          20 BP; 4 A;
                               AGCTGGGACTACAGGCGCCC 751
AGCTGTGACTACAGGTGCCC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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                                                                                                                          6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                           90.0%;
                                                                                          1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 16.8; DB 1;
Pred. No. 1.6e+03;
                                                                           Score 16.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                               Mismatches
                                                                             1.6e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                              DB 1;
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                                                                                            Length 20;
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                                                               Indels
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                                                              0
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                                                               Gaps
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RESULT 1336
ABL4402
ID ABL4402
XX
AC ABL4402
XX
AC ABL4402
XX
DT 11-APR-

ABL44022;

ABL44022 standard; DNA; 20

В₽

11-APR-2002

(first entry)

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RESULT 1335
ABL4433
AC ABL4433
XX ABL4433
XX ABL4433
XX Human C
XX Human C
XX Human C
XX Human C
XX Homo Ba
XX PCR Pri
XX PR 10-MAR-
XX (RIKA)
PA (RIK
                                                                                                                                           Matches
                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                               plates; (e) the clones in the multiwell plates of the specified discrimination Nos. are mixed respectively in each wells of longitudinal and lateral directions; (f) the mixed clones are cultured and the resultant cultures are amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABL42957 to ABL45322 represent PCR primers for human chromosome 1936-35 DNA, and ABL45323 to ABL45634 represent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell the maximum in the specified discrimination Nos.
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                                                                                                                                                                                                                                                                                   Sequence 20 BP; 4 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Arraying genome clones.
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                                                                                                                                                                             1.7%;
Similarity 90.0%;
CTCACTGCAGCATTCACCTC 20
                                                                                                                                           Conservative
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                                                                                                                                                                             Score 16.8; DB 1;
Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                          0 Other;
                                                                                                                                                                                                               Length 20;
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Human chromosome 1p36-35 PCR primer SEQ ID NO:1066

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ABL44316
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and lateral directions; (f) the mixed clones are cultured and the resultant cultures are amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABI42957 to ABI45323 represent PCR primers for human chromosome 1p36-35 DNA, and ABI45323 to ABI45634 represent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention
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                Homo sapiens
                                                                                                        Human chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 5 A; 7 C; 3 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 26; 528pp; Japanese
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                                                                   Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
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                                                PCR primer;
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                                                                                                                                                                                                                                                                                                                             379 TCAGCCTCCCAAAGTGCTGG 398
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                                                                                                                                          (first entry)
                                                                                                        1p36-35 PCR
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                                                                                                      primer SEQ ID NO:1360
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Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                1.6e+03
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ARKSULT 1338
ABK68938
ID ABK6893
AC ABK6893
XX ABK6893
XX DT 02-JUL-
XX DE Human p
XX Human,
KW infecti
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KW antiinf
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Best Local Similarity
                                                                                                                                                                                              Human; phosphorylase kinase beta; metabolic disorder; diabetes; infection; inflammation; tumour formation; antidiabetic; antiinflammatory; cytostatic; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 4 A; 9 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; Page 31; 528pp; Japanese
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-MAR-2001; 2001JP-00068285
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                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                Human phosphorylase kinase beta antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                        02-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                      ABK68938 standard;
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                                                                                                                                                                                                                                                                                                                                                               ABK68938;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         specifically claimed
                                                                                                   modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          931 CTCACTCTGTTACCCAGGCT 950
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primers for human chromosome 21q22.1, lly claimed for use in the present inventi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                        (first entry)
/mod_base= OTHER
/note= "OTHER= Phosphorothioate internucleotide linkages,
optionally bases 1-5 and 16-20 are 2'-methoxyethoxy (2'-
MOE) bases, where the 2'-MOE cytidines are also
                                                                              /*tag=
                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
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Pred. No. 1.6e+03
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Same and a state of

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RESULT 1339
AAL38209
ID AAL38209;
XX AAL38209;
XC AAL38209;
XY 15-AUG-20
DT 29-AUG-20
XX 15-AUG-20
XX Hepatotro
KW haemostat
KW haemostat
KW haemostat
KW haemostat
KW 2'-MOE; p
XX
PM HOMO sapi
OS Chimeric.
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PD 114 MAR-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to antisense compounds and methods for compoulating the expression of human phosphorylase kinase beta. The antisense compounds, particularly antisense oligonucleotides, target and cinhibit the expression of human phosphorylase kinase beta. The antisense compounds are useful for inhibiting the expression of human phosphorylase kinase beta. The antisense compounds are useful for inhibiting the expression of human phosphorylase kinase beta in human cells or tissues and for treating an animal, comparticularly a human suspected of having or being prome to a disease or condition associated with expression of phosphorylase kinase beta such as a metabolic disorder e.g. diabetes. The compounds are useful for compartice, therapeutics and as research reagent, e.g. prophylactically to prevent or delay infection, inflammation or tumour formation. The compounds are useful in the preparation of a pharmaceutical formulation. They are highly specific, have an enhanced affinity for the nucleic acid target, and are safely and effectively administered to humans. ABK6888-ABK68965 represent human phosphorylase kinase beta compounds which comprise a phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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                                                                                                                 haemostatic; BH3 interacting domain death agonist; liver disease; haematopoietic disorder; developmental disorder; immunological disorder; hyperproliferative disorder; apoptosis; human; chimeric; 2'-methoxyethyl; 2'-MOE; phosphorothicate backbone; ds.
                                                                                                                                                                                                                                                                                                           AAL38209;
                                                                                                                                                                                                                                                                                                                                          AAL38209 standard;
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                                                                                                                                                                                      Hepatotrophic; immunomodulatory;
                                                                                                                                                                                                                      Human BH3 interacting domain death mRNA agonist inhibitor SEQ
                                                                                                                                                                                                                                                         29-AUG-2003
15-AUG-2002
                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                             675
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                                                                                                                                                                                      cytostatic; antiinflammatory; hepatitis;
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14-MAR-2002.

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The invention relates to a compound 8 to 50 nucleotides in length CC targeted to a nucleic acid molecule encoding a BH3 interacting domain CC death agonist, where the compound specifically hybridises with and CC inhibits the expression of the BH3 interacting domain death agonist. The Cc compound of the invention is useful for inhibiting the expression of the CC also useful for treating an animal having a disease or condition CC associated with the BH3 interacting domain death agonist, e.g. CC haematopoietic disorder, hyperproliferative disorder, a developmental CC disorder, immunological disorder, or a disease or condition of the liver Cc., hepatitis, or a condition associated with apoptosis. The compound CC is useful for disgnostics, therapeutics, prophylaxis and as research CC easy, hepatitis, or a condition associated with apoptosis. The compound CC reagents and kits. This polymucleotide sequence represents an antisense CC oligonucleotide inhibitor of the DNA from human BH3 interacting domain CC death agonist RNA of the invention. NOTE: This sequence is a chimeric CC plosphorotides. The internucleoside (backbone) linkages are phosphorothioate (P=S) throughout the oligonucleotide. (Updated on 29-AUG cc. 2003 to standardise OS field)
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel antisense compound targeted to nucleic acid molecule encoding BH3 interacting domain death agonist, useful for treating animals we diseases associated with BH3 interacting domain death agonist, e.g.
                                                                                                          Human; PCR; primer; ornithine decarboxylase; odc; 88; 8usceptibility; epithelial cancer; A-allele; G-allele; polyamine level; carcinogenesi single nucleotide polymorphism; SNP; molecular baccon probe; skin; digestive system; oesophageal; gastric; colon; prostate; breast; haematopojetic; lung; cervical; cancer; melanoma; carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-AUG-2001; 2001WO-US027316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 3; Page 87; 171pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-393838/42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                      772
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                            TTGTATTTTTAGTAGAGATG 791
                                                                                                                                                                                                                                                                                                                                                                                        TTGTATTTTAAGTAGAGACG 20
                                                                                                                                                                                                         primer 1, for allelic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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2001US-00800631.
                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 C; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90.0%;
                                                                                                                                                                                                                                                                                                            20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 16.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                              discrimination of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ŏ.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              acid molecule encoding the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                odc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20;
                                                                                                                                                                 carcinogenesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               animals with
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                                                                                                                                                                                                                -3175
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                                                                                                                                                                                                                SNP.
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RESULT 1341
ABK11979
ID ABK1197
XX ABK1197
XC ABK1197
X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC determining whether the human comprises an A-allele of the ornithine decarboxylase (odc) gene, where its presence indicates a greater cc decarboxylase (odc) gene, where its presence indicates a greater cc susceptibility to epithelial cancer than one without the allele. Odc is involved in establishing cellular polyamine levels and the susceptibility cof a tissue to carcinogenesis is related to these polyamine levels. This cc an be achieved by determining the sequence of a region of the gene cc containing the single nucleotide polymorphism (SNP) or by contacting a cc polynucleotide derived from the human's genome with a first molecular cc beacon probe which is complementary to a SNP target region of the odc cg gene (e.g. at positions -3175, -3004, -1936, +263, +317, +5294, +5915, +6697, +7487 or +7886 relative to the transcription start site of the cc gene). The invention discloses a kit for assessing susceptibility of a human to an epithelial cancer which comprises the primer and coligonucleotide probes for determining the presence or absence of the A-cc allele. The method is useful in assessing the relative susceptibility of a human to an epithelial cancer, such as skin, digestive system, componed is an inhibitor or an inducer of carcinogenesis. The sequence cc presented is the forward PCR primer 1, which was used for allelic concernination of the human odc -3175 SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Assessing susceptibility of humans to epithelial cancer comprises the determination of A- or G-allelles of the ornithine decarboxylase (odc) gene which is an indicator of susceptibility to epithelial cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-MAR-2000; 2000US-00516357
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      17-JUL-2001; 2001US-00907190.
                                                                                                                                                                                                                                                                                            Human; linkage; familial dysautonomia; FD; D9S58; rchromosome 9q31-q33; prenatal diagnosis; Riley-Day
                                                                                                                                                                                                                                                                                                                                                                                                Human D9S58
                                                                      28-FEB-2002
                                                                                                                                     US2002025528-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABK11979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention discloses a method for assessing the relative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (LANK-) LANKENAU MEDICAL RES CENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABK11979 standard; DNA; 20
                                                                                                                                                                                                 sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
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                                                                                                                                                                                                                                                                                                                                                                                             genetic marker PCR Primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                             D9S58; neuronal loss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                               syndrome;
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RESULT 1342
ABS65069/c
ID ABS6506
XX
AC ABS6506
XX
DT 15-NOV-
XX
DE Human C
XX
KW Ss; ant
KW Cytosta
KW Cytosta
KW hyperpx
KW 1iver C
XX
Hömo sa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to a novel method for detecting a polymorphism CC linked to a gene associated with familial dysautonomia (FD). Familial CC dysautonomia is an autosomal recessive disorder characterised by the CC developmental loss of neurons from the sensory and autonomic nervous CC system. The method of the invention comprises analysing human chromosome CC and detecting the presence of a polymorphism located between the CC genetic markers 19853 and 1983105 inclusive, and linked to the gene CC associated with familial dysautonomia. The invention also includes CC nucleotide sequences for detecting a polymorphism associated with CC familial dysautonomia. Using the method of the invention it was possible CC show that the gene for FD is located on human chromosome 9q31-q33. The CC method and sequences of the invention are useful for the diagnosis of CC method and sequences of the invention are useful for the diagnosis and CC disease gene, such information will facilitate prenatal diagnosis and CC help reduce the number of new cases of FD. The present sequences CC represent an oligonucleotide primer that can be used to screen for the CC 19858 genetic marker on chromosome 9, this primer was used to map the CC location of the familial dysautonomia gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-MAY-1992;
16-APR-1993;
07-JUN-1995;
                                                                                                                               cytostatic; antidia hyperproliferative liver cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 5 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 6; 17pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detecting a polymorphism linked to a gene associated dysautonomia, involves analyzing human chromosome 9 f
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Blumenfeld A, Gusella
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(GUSE/)
(BREA/)
                                                                                                                                                                                                                Human casein
                                                                                                                                                                                                                                                 15-NOV-2002
                                                                                                                                                                                                                                                                                 ABS65069;
                                                modified_base
                                                                                                 Homo sapiens.
                                                                                                                                                                                ss; antisense;
                                                                                                                                                                                                                                                                                                             ABS65069 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                            725 CCTGAGTAGCTGGGACTACA 744
                                                                                                                                                                                                                                                                                                                                                                                                                                                            18;
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BREAKFIELD X
SLAUGENHAUPT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                              CCTGAGTAGCCGGGACTATA 20
                                                                                                                                                se; casein kinase2-beta; human; antisense gene therapy;
antidiabetic; antiinflammatory; diabetes; cancer; tumour;
erative disorder; breast cancer; prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                 kinase
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93US-00049678.
95US-00480655.
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                                                  Location/Qualifiers
1. .20
                /*tag=
/mod_ba
/mod_base= OTHER
/note= "All cytidines are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                 2-beta antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 20;
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                                                                                                                             Query Match
Best Local
                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            kinase 2-beta, useful
treating a disease or
kinase 2-beta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides targeted to nucleic acid encoding Casein kinase 2-beta, useful in diagnostic and research applications, or for treating a disease or condition associated with the expression of Casein
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                                                                                                      l Similarity
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                                     CTCGGCTCACTGCAACCTCT 988
     CTCGGCTTACTGCCACCTCT 1
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                                                                                                         Conservative
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16. .20
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/note= "2'-methoxyethyl residues"
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                *tag=
                                                                                                                                                                                                                Ą
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             _base= OTHER
e= "2'-methoxyethyl residues"
                                                                                                                                   90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wyatt JR;
                                                                                                                                                                                                                C; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Phophorothioate
                                                                                                      0;
                                                                                                                                                                                                                   G; 2 T;
                                                                                                                                      Score 16.8;
Pred. No. 1
                                                                                                         Mismatches
                                                                                                                                                                                                                   o
u;
                                                                                                                                      1.6e+03;
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                                                                                                                                                                                                                   0
                                                                                                                                                            DB 1;
                                                                                                                                                                                                                   Other;
                                                                                                                                                          Length 20;
                                                                                                            Indels
                                                                                                         0
                                                                                                            Gaps
                                                                                                         0;
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RESULT 1343
AAL45049
ID AAL4504
XX
AC AAL4504

standard;

DNA;

20

AAL45049; AAL45049

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         airway hypersensitivity; autoimmune disease; endotoxin shock; sepsis; microbial infection; hepatitis B; hepatitis C; diabetes; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hepatitis C, insulin-weptimed diseases. The present sequence wrotein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention provides the protein and coding sequences of several protein capable of enhancing the activity of NF-kappaB. These be used in the treatment of allergy, atrophy, asthma, pollenosis, airwhypersensitivity, autoimmune diseases, graft-vs.-host diseases, endoto shock, sepsis, microbial infections, chronic hepatitis B, chronic hepatitis C, insulin-dependent or independent diabetes and many other diseases. The present sequence is a PCR primer for a coding sequence of the sequence of the primer for a coding sequence of the sequence of the present sequence is a PCR primer for a coding sequence of the sequence of the present sequence of the present sequence of the present sequence of the primer for a coding sequence of the present sequence of the presen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A new polypeptide useful in the development of agents to treat {\bf e}. autoimmune diseases and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP2001352986-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human NF-kappaB activity enhancer PCR primer
                                                                                                                                                                                                                                                               Human; cyclin-dependent kinase; CDK; cyclin-dependent kinase inhibitor; inhibitor; cancer; age-related disease; promoter; atherosclerosis; cytostatic; antiarteriosclerotic; nootropic; neuroprotective; nephrotropic; antiarthritic; arthritis; renal disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-191857/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-MAY-2002 (first entry)
01-FEB-2001; 2001US-0265840P
                                                                                                                                                                                                                                                                                                                                                                                        Human cathepsin B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABT08416 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 4; Page 49; 52pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (KYOW ) KYOWA HAKKO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-JUN-2000; 2000JP-00175475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-JUN-2000; 2000JP-00175475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                01-FEB-2002; 2002WO-US002784.
                                                                                                                                                WO200266681-A2
                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                             Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                        27-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABT08416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 4 A; 7 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                      promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KOGYO KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВP
                                                                                                                                                                                                                                                                                                                                                                                        PCR primer SEQ ID NO: 51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. 1.6
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #11.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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RESULT 1345
AAD55399
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a recombinant expression construct encoding a reporter gene operably linked to a promoter from a mammalian gene induced by a cyclin-dependent kinase (CDK) inhibitor. The construct is useful for identifying compounds that inhibit the induction of genes induced by CDK inhibitors. The compounds are useful for preventing or treating a disease caused by CDK inhibitor induced gene expression, e.g. cancer other than colon cancer, renal failure, Alzheimer's disease, amyloidosis, age-related diseases, atherosclerosis or arthritis. The present sequence is a PCR primer used to amplify a human promoter suitable for use in the construct of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New recombinant expression construct, useful for identifying compounds that inhibit the induction of genes induced by cyclin-dependent kinase inhibitors for preventing or treating cancer, renal failure or Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                interferon-induced double stranded RNA-activated p68 kip1/eIF2 alpha protein kinase; gene therapy; infection; phosphorothioate backbone; ss.
                                                                                                                                                                                                                                                                                                                                             Human; protein kinase R; PKR; PRKR; immunosuppressive; antiinflammatory; interferon-induced double stranded RNA-activated p68 kinase; DAI; dsI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAY-2001; 2001US-00861925
                                                                                                                                                                                                                                                                         Homo sapiens
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                          Human PKR antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                       07-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAD55399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 8; Page 129; 137pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UNII ) UNIV ILLINOIS FOUND.
20-MAR-2003.
                           WO2003022222-A2
                                                                                                modified_base
                                                                                                                                                     modified_base
                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAD55399 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002-674960/72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       723 CTCCTGAGTAGCTGGGACTA 742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       μ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Roninson IB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                     /*tag= b
/mod_base= OTHER
                                                                                                                                                                    /note= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
                                                                                                                                                                                                              /*tag=
                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                              note=
                                                                                                                                                                                                  mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.7%;
                                                                     _base= OTHER
                                                      "2'methoxyethyl nucleotides"
                                                                                                               "2'methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 16.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8
                                                                                                                                                                                                                                                                                                                                                                                          ISIS 139452.
                                                                                                                 nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.6e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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Matches 18
                                                                                                                                                                                   dal, and PI/eIF2 alpha protein kinase). The compositions contain antisense compounds, particularly antisense oligonucleotides targetted t nucleic acids encoding PKR. The antisense compound is useful for inhibiting the expression of PKR and for modulating the process of RNA-mediated interference (RNAi) in a cell. It is useful for treating an animal having a disease or condition associated with PKR. It is also useful for distinguishing functions of various members of biological and kits, for distinguishing functions of various members of biological pathway, and in antisense gene therapy. It is useful prophylactically, e.g., to prevent or delay infection, inflammation or tumour formation. The present sequence is an antisense oligonucleotide targetted to human PKR DNA. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel antisense compound that hybridizes and inhibits nucleic ac encoding protein kinase R, useful for treating animal having dis condition associated with protein kinase R such as an autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ward DT,
                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to antisense compounds, compositions and methods for modulating the expression of protein kinase R (also known as PKR, pRKR, interferon-induced double stranded RNA-activated p68 kinase, DAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 15; Page 77; 61pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-SEP-2002; 2002WO-US028870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM
                                                                          Local Similarity
les 18; Conserv
                          220 AACTCCCGACCTCAGATGAT 239
  _
                                                                                                                                                      20 BP; 5 A; 6 C; 4 G; 5 T; 0 U; 0 Other;
  AACTCCTGACCTCAGGTGAT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Watt AT;
                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2001US-00953611
                                                                                            90.0%;
                                                                          <u>.</u>
                                                                                            Score 16.8;
Pred. No. 1.
                                                                            Mismatches
                                                                                                1.6e+03
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                                                                                                                Length
                                                                              Indels
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RESULT 1346
ABZ79344
ID ABZ7934
WO2002100896-A2
                                                                                                                                    Acetyl-Coenzyme A-carboxylase-alpha gene PCR primer, SEQ ID
                                                                                                                                                  01-MAY-2003
                                                                                                                                                                ABZ79344;
                                                                                                                                                                              ABZ79344 standard; DNA; 20
                                                                                                 Homo sapiens
                                                                                                                     Human; enzyme;
                                                                                                                                                  (first entry)
                                                                                                              acetyl-Coenzyme A-carboxylase-alpha; ACC-alpha; cancer;
PCR; primer; ss.
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Dalla Venezia NL,

Magnard CM,

Lenoir GM,

Sinilnikova-Erard

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(CNRS ) CNRS (UYLY-) UNIV

CENT NAT RECH SCI. LYON 1 BERNARD CLAUDE.

13-JUN-2001; 05-MAR-2002;

2001FR-00007740. 2002FR-00002788.

12-JUN-2002; 2002WO-FR002015.

19-DEC-2002

2003-175165/17

breast and c

l ovarian me acetyl

coenzyme cancer,

0

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RESULT 1347
ACC79697/c
ID ACC7969
XX ACC7969
XX ACC7969
XX ACC7969
XX ACC7969
XX Human;
KW Human;
KW Apoptos
KW Chemoth
XX Homo sa
OS Synthet
XX WO2003C
XX ACCACA
PN WO2003C
XX Homo sa
OS Synthet
XX Human;
KW Apoptos
XX ACCACA
PN WO2003C
XX Human;
PF 17-SEP-
XX EIRX-)
PA (EIRX-)
PA (EI
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              alpha (ACC-alpha; see ABZ79442), which can be used for in vitro diagnosis of cancer (or of an increased risk of developing it), by detecting ACC-alpha gene mutations or polymorphisms, or altered ACC-alpha protein expression, relative to a control population. The method is particularly used to diagnose cancer, especially of breast or ovary, or for assessing the risk of developing such cancers. The present sequence is a PCR primer, which was used in an example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              In vitro diagnosis of cancer, particularly break susceptibility, comprises detecting alterations carboxylase alpha gene or protein expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; DELTA-N p73; apoptosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACC79697 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chemotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACC79697;
                                                                                                                                                                                                     New human delta-N p73 proteins and nucleic acids encoding them, usefu
for diagnosing, preventing and treating diseases associated with
decreased or increased apoptosis, or for predicting a predisposition
                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-SEP-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                   The present invention describes isolated human DELTA-N p73
                                                                                                                                      Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003025010-A2
  molecules (I). (I) have apoptotic and anti-apoptotic activities, and cobe used in protein therapy. The DELTA-N p73 nucleic acids may be used inhibiting apoptosis or the expression of a p53, p63, or an N-terminal transactivation (TA) p73 molecule in a cell, for predicting tumour
                                                                                                                                                                                                                                                                                                                                                                                                          (EIRX-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       382
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                                                                                                                                                                                                                                                                                                                                                                                                             EIRX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                     Melino G, I
a F, Tobler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 10; 56pp; French.
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                                                                                                                                                                                                                                                                                                                                                                                          uman; mouse; ss; primer; gene 216; antiasthmatic; antiinflammatory; norectic; chromosome 20p13-p12; single nucleotide polymorphism; SNP; ene therapy; respiratory disease; asthma; obesity; PCR; ronchial hyper-responsiveness; chronic obstructive pulmonary disease;
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This invention relates to a novel isolated nucleic acid, gene 216, identified from human chromosome 20p13-p12. The invention also discloses regions of the 216 gene that contain single nuclectide polymorphisms (SNP's) which may be used as markers for disease susceptibility or severity. The nucleotides of the invention may have antiasthmatic, antiinflammatory or anorectic activities and may be used in gene therapy. The nucleic acids, antibodies or its fragments are useful for diagnosing, preventing or treating a disorder, such as respiratory diseases (e.g. asthma, bronchial hyper-responsiveness, chronic obstructive pulmonary disease or adult respiratory distress syndrome), obesity, or inflammatory

bowel syndrome.

The nucleic acids

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RESULT 1349
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                                                                                                                                                                                                                                                                                     New antisense oligonucleotide targeted to a nucleic acid encoding vascular endothelial growth factor receptor-1, useful for diagnosing treating cancer, rheumatoid arthritis, or diseases or conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vascular endothelial growth factor receptor 1; VBGF receptor; VBGFR; inhibitor; cytostatic; antirheumatic; antiarthritic; antiangiogenic; antiinflammatory; antisense gene therapy; hyperproliferative disorder; cancer; rheumatoid arthritis; angiogenesis; infection; inflammation; tumour formation; phosphorothioate; 2'-O-methoxyethyl; 2'-MOE; ss.
                             targeted to a nucleic acid molecule encoding vascular endothelial growth factor receptor-1 (VBGFR-1), where the compound inhibits the expression of VEGFR-1 and specifically hybridises with the nucleic acid encoding VEGFR-1 or with an 8-nucleobase portion of an active site on the nucleic
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                                                                                                                                              The present invention describes a compound (C) 8-50 nucleobases in length
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  8-nucleobase portion of an active site on ding VEGFR-1. Also described: (1) a composi
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CC comprising (C) and a carrier or diluent; (2) inhibiting the expression of VEGFR-1 in cells or tissues by contacting the cells or tissues with (C) CC so that the expression of VEGFR-1 is inhibited; and (3) treating an CC administering (C) to the animal so that the expression of VEGFR-1 by CC administering (C) to the animal so that the expression of VEGFR-1 is CC inhibited. (C) has antiangiogenic, antirheumatic, antiarthritic, CC cytostatic and antiinflammatory activities, and can be used in antisense compounds are useful for modulating the CC expression of VEGFR-1 and for treating diseases or conditions associated CC with the expression of VEGFR-1, such as hyperproliferative disorders (c.e.g. cancer), rheumatoid arthritis, or diseases or conditions involving CC angiogenesis. The antisense compounds are also useful for diagnostics, CC therapeutics, prophylaxis, e.g. to prevent or delay infection, and in CC distinguishing between functions of various members of a biological CC pathway. The present sequence represents a human VEGFR-2 chimeric phosphorothicate antisense coligonucleotide, which is used in an example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; HKR1; cytostatic; HKR1 inhibitor; hyperproliferative disorder; cancer; antisense oligonucleotide; 2'-O-methoxyethyl; 2'-MOE; control phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABZ71057 standard;
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                                                 03-JUL-2001;
                                                                                                   02-JUL-2002; 2002WO-US021090
                                                                                                                                                                                                                                                                                                                              modified_base
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                                                                                                                                                    16-JAN-2003
                                                                                                                                                                                                      WO2003004513-A1.
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  PHARM INC.
                                                 2001US-00898556
                                                                                                                                                                                                                                                                                                                              /note= "
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag=
/mod_ba
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                                                                                                                                                                                                                                                                                  mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         'note= "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                 base= OTHER
e= "2'-O-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   base= OTHER
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Pred. No. 1.
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                                                                                                                                                                                                                                                        nucleotides"
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WPI; 2003-210336/20

Bennett FC,

Freier

Example 15; Page

73; 105pp; English.

to,

invention describes a compound 8-50 nucleobases in length and which specifically hybridises with a nucleic acid

associated with nucleic

acid

s, particularly antisense oligonucleotides targeted to a encoding HKR1, useful for treating a disease/condition ith HKR1, such as hyperproliferative disorder, e.g. lung.

such as hyperproliferative

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RESULT 1351
ABZ71059/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC molecule encoding HRR1, and inhibits the expression of HRR1. Also composited: (1) a compound 8-50 nucleobases in length that specifically composited and at least an 8-nucleobases in length that specifically composited acid molecule encoding HRR1; (2) a composition of an active site on a cc nucleic acid molecule encoding HRR1; (2) a method for inhibiting the cc compound and a carrier or diluent; (3) a method for inhibiting the cc expression of HRR1 in cells or tissues by contacting the cells or tissues with the compound so that expression of HRR1 is inhibited, and (4) a cc method of treating an animal having a disease or condition associated with HRR1 by administering to the animal a therapeutic or prophylactic amount of the compound so that expression of HRR1 is inhibited. HRR1 can be used as the string a disease or condition and methods are useful for treating a disease or comdition and methods are useful for treating a disease or condition associated with HRR1, such as a cc hyperproliferative disorder, e.g. lung, brain or breast cancer, by diagnostics for modulating the expression of HRR1. They are also useful in research and creating the expression of HRR1. They are also useful in research and conjugations of the system of the present sequence conjugated useful in the inhibition of human HRR1 in an antisense coligonucleotide used in the inhibition of human HRR1 in an example from
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                     Key
modified_base
                                                                                                                                                                                                                                                                               Human; HKR1; cytostatic; HKR1 inhibitor; hyperproliferative disorder;
cancer; antisense oligonucleotide; 2'-O-methoxyethyl; 2'-MOE; control;
                                                                                                                                                                                                                                                                 phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                                    ABZ71059
                                                                                                                                 modified_base
                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                            Human
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 WO2003004513-A1
                                                                      modified_base
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                                                                                                                                                                                                                                                                                                                            HKR1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                     standard; DNA;
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                                                                                                                                                                                                                                                                                                                         phosphorothicate antisense oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention
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                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                            note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.7%;
                                                                                     base=
e= "2'
                             base= OTHER
== "2'-O-methoxyethyl
                                                                                                                                                             base=
                                                                                                                                               "phosphorothioate linkages"
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                                                                                                                                                                                                                                                                                                                                                                                                                     20
                                                                                       -O-methoxyethyl
                                                                                                                                                             OTHER
                                                                                                     OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>,,</u>
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Pred. No. 1.6e+03;
0; Mismatches 2
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                               (2'-MOE)
                                                                                       (2'-MOE)
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                               nucleotides'
                                                                                       nucleotides"
                                                                                                                                                                                                                                                                                                                               SEQ
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The present invention describes a compound 8-50 nucleobases in length cargeted to, and which specifically hybridises with a nucleic acid compound encoding HKR1, and inhibits the expression of HKR1. Also ceasing the compound s-50 nucleobases in length that specifically compounds and a carrier or diluent; (3) a method for inhibiting the compound and a carrier or diluent; (3) a method for inhibiting the compound and a carrier or diluent; (3) a method for inhibiting the compound and a carrier or diluent; (3) a method for inhibiting the compound and a carrier or diluent; (3) a method for inhibiting the compound so that expression of HKR1 is inhibited; and (4) a compound of treating an animal having a disease or condition associated compount of the compound so that expression of HKR1 is inhibited; and (4) a compount of the compound, composition at the appeutic or prophylactic compount of the compound, composition and methods are useful for treating a disease or condition associated with HKR1 such as a compound; composition and methods are useful for the special compound of HKR1. They are also useful in research and composition for modulating the expression of HKR1. They are also useful in research and composition be used as human HKR1 chimeric phosphorothicate of present sequence conjugonucleotide used in the inhibition of human HKR1 in an example from the present sequence conjugonucleotide used in the inhibition of human HKR1 in an example from the present is an antisense conjugonucleotide used in the inhibition of human HKR1 in an example from the present is an antisense conjugonucleotide with the inhibition of human HKR1 in an example from the present is an antisense conjugonucleotide with the inhibition of human HKR1 in an example from the present is an antisense conjugonucleotide with the inhibition of human HKR1 in an example from the present is an antisense conjugonucleotide with the present is an antisense conjugonucleotide with the present is an antisense conjugonucleotide with the present is an antisense co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New compounds, particularly antisense oligonuclectides targeted to nucleic acid encoding HKR1, useful for treating a disease/condition associated with HKR1, such as hyperproliferative disorder, e.g. lum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-210336/20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            brain or breast cancer.
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문 S Query Match Best Local 9 Matches Sequence present invention 930 20 18; Similarity 20 TCTCACTCTGTTACCCAGGC 949 BP; 7 A; 4 C; 7 Conservative 90.0%; G; 2 T; 0 U; 0 Other; 0; Score 16.8; DB 1; Pred. No. 1.6e+03 Mismatches 2 Length 20; Indels 0 Gaps

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RESULT 1352
ADA20923/c
                                                                                                BCL2-associated X; BAX; nootropic; neuroprotective; antipanticonvulsant; ophthalmological; antidiabetic; virucide;
                                                                                                                                             Human BAX
                                                                                                                                                                           20-NOV-2003
                                                                                                                                                                                                         ADA20923;
                                                                                                                                                                                                                                     ADA20923
                        Parkinson's disease; Hodgkin's disease; cartilage-hair hyperplasia;
diabetes-associated ocular disorder; scrapie infection;
aberrant apoptosis; human; phosphorothioate; ss.
                                                                       amilianse therapy; BAX antagonist; BAX inhibitor; amilial amylotrophic lateral sclerosis; Alzheimer's disease;
                                                                                                                                                                                                                                      standard; DNA;
                                                                                                                                               chimeric
                                                                                                                                                                           (first entry)
                                                                                                                                            phosphorothicate oligonuclectide
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                                                                                                                                                                                                                                      ₽₽
                                                                                                                   antiparkinsonian;
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                       δ
                                                                                                                                                                                                       crangeted to a nucleic acid molecule encoding BCL2-associated X (BAX)

composed protein, where the compound specifically hybridises with the nucleic acid molecule encoding BAX protein and inhibits the expression of BAX protein. The compound specifically hybridises with at least 8-nucleobase portion of an active site on a nucleic acid molecule encoding BAX protein. Also described: (1) a composition comprising (I) and a pharmaceutical carrier or diluent; (2) inhibiting the expression of BAX protein in cells or tissues comprising contacting the cells or tissues with (I); and (3) treating an animal having a disease or condition associated with BAX protein is inhibited. (I) has nootropic, neuroprotective, antiparkinsonian, anticonvulsant, ophthalmological, antidiabetic and colinarist The antisense compounds (I) are useful for modulating the expression of BAX protein, and for treating a disease or condition associated with BAX protein, e.g. familial amylotrophic lateral sclerosis, Alzheimer's disease, Parkinson's disease, Hodgkin's disease, cartilage-hair hyperplasia, diabetes-associated ocular disorders or contition or a condition that arrises from abarrant approach.
                                                           Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compounds, useful for modulating the expression of BCL2-associated X (BAX) protein or for treating a disease or condition associated with BAX protein, e.g. Parkinson's disease, Hodgkin's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo
                                                                                                                                               scrapie infection, or a condition that arises from aberrant apoptosis. The compounds are useful as research reagents and in diagnostics. The present sequence represents a human BAX chimeric phosphorothicate oligonucleotide, which is used in an example from the present invention
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                                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3; Page 87; 139pp;
20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ISIS PHARM INC
                                                                         Similarity
                            GGAGTGCAGTGGCGCAATCT
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GGAGTGCAATGGCGCAACCT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention describes a compound (I) 8-50 nucleobases in length
                                                                                                                    B₽;
                                                           Conservative
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16. .20
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                                                                      1.7%;
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); Mismatches
                                                                       Score 16.8;
Pred. No. 1.
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                                                                                                                                                     invention
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RESULT 1353
ADA20924/c
        The present invention describes a compound (I) 8-50 nucleobases in length (I targeted to a nucleic acid molecule encoding BCL2-associated X (BAX) (I protein, where the compound specifically hybridises with the nucleic acid molecule encoding BAX protein and inhibits the expression of BAX protein. The compound specifically hybridises with at least 8-nucleobase portion (I a nactive site on a nucleic acid molecule encoding BAX protein. Also (I a composition comprising (I) and a pharmaceutical carrier or diluent; (2) inhibiting the expression of BAX protein in cells or tissues comprising contacting the expression of BAX protein in cells or tissues comprising analimal having a disease or condition associated with BAX (I reating an animal having a disease or condition associated with BAX (I protein comprising administering to the animal (I) so that expression of BAX protein is inhibited. (I) has nootropic, neuroprotective, antiparkinsonian, anticonvulsant, ophthalmological, antidiabetic and contributes, and can be used in antisense therapy, and as a BAX antigenist. The antisense compounds (I) are useful for modulating the
                                                                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA20924 standard;
                                                                                                                                                                                                                                                                                                                  New antisense compounds, useful for modulating the expression of BCL2-associated X (BAX) protein or for treating a disease or condition associated with BAX protein, e.g. Parkinson's disease, Hodgkin's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Parkinson's disease; Hodgkin's disease; cartilage-hair hyperplasia; diabetes-associated ocular disorder; scrapie infection; aberrant apoptosis; human; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human BAX chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADA20924;
                                                                                                                                                                                                                                                                                                        or Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-JUL-2001; 2001US-00908147.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-JAN-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense therapy; BAX antagonist; BAX inhibitor; familial amylotrophic lateral sclerosis; Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BCL2-associated X;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-NOV-2003
                                                                                                                                                                                                                                                                         Claim
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                                                                                                                                                                                                                                                                     Page 87; 139pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                    Watt AT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "phosphorothioate linkages, residues are 5-methylcytidines"
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_e= "2'.
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RESULT 1354
ADB16959
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cartilage-hair hyperplasia, diabetes-associated ocular disorders or scrapie infection, or a condition that arises from aberrant apoptosis. The compounds are useful as research reagents and in diagnostics. The present sequence represents a human BAX Chimeric phosphorothicate oligonucleotide, which is used in an example from the present inventic
                             This invention relates to a novel isolated human gene DYXC1 that is functionally related to dyslexia, more particularly it describes single nucleotide polymorphisms thought to predispose an individual in to developing dyslexia. This is a neurological disorder with a genetic basis (DYXC1 has been isolated to chromosome 15q21), which manifests itself as a specific reading disability. Specifically, DYXC1 is can be useful in study of brain processes such as reading, phonological processing, rapid naming and verbal short-term memory. Accordingly, the present invention describes methods and materials for analysing allelic variations in the DYXC1 gene, and also provides DYXC1 as an antigen for the production of antibodies used in the diagnosis of dyslexia. This oligonuclectide is the EKN1-9R PCR primer that is specific for human intronic DYXC1, and is used to amplify exon 9 in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 associated with BAX protein, e.g. familial amylotrophic lateral sclerosis, Alzheimer's disease, Parkinson's disease, Hodgkin's disease,
                                                                                                                                                                                                                                                                                                                                      New isolated, purified DYXC1 nucleic acid for studying brain processes, e.g. reading, phonological processing, rapid naming or verbal short-term memory, or for diagnosing dyslexia or assessing the predisposition to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo
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                                                                                                                                                                                                                                                                                   Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LICENTIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disability; phonological processing; rapid naming;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                   Page 23; 135pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LTD.
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                                                                                                                                                                                                                                                                                     English.
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Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer
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Query Match

Sequence

20

BP;

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A; 5 C;

5 G;

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0 Other;

RESULT 1356 ABT44385/c ID ABT4438 XX AC ABT4438

standard; DNA;

20

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ABT44385 ABT44385

1.7%;

Score 16.8; 6 T;

DB 1;

Length

20;

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RESULT 1355
ADA27318/c
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Matches 18; Conserv
                                                               Query Match
Best Local (
                                                   Matches
                                                                                                                                                      The invention relates to an oligonucleotide primer capable of specifically hybridising to a DNA having the sequence of the flanking regions of a microsatellite selected from M2-4-9, M2-2-12, M2-2-12, M2-2-21, M2-2-22, M2-2-23, M2-2-24, M2-2-24, M2-2-25, M2-4-36, M2-2-26, M2-2-36, M2-4-31, M2-3-22, M2-2-36, M2-5-11, M2-2-29, M2-2-32, M2-4-32, M2-4-33, M2-4-37, M2-3-22, M2-2-36, M2-5-11, M2-2-46, and M2-2-48. The primer is useful for determining the number of repeat units of the microsatellite cited above. The primer is useful in HIA-related research, such as genetic mapping of HIA class II-associated diseases, transplantation matching, population genetics, and identification of recombination hot spots as well as linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADA27318
                                                                                                                             disequilibrium studies. The present sequence microsatellite M2_4_9 PCR primer \#2.
                                                                                                                                                                                                                                                                                                            Claim 4; Page 7; 20pp; English.
                                                                                                                                                                                                                                                                                                                                      New oligonucleotide primer capable of specifically hybridizing to a DNA having the sequence of the flanking regions of a microsatellite (e.g. M249), useful for HLA-related research, e.g. transplantation matching.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-DEC-2002; 2002US-00314405.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ss; primer; PCR; HLA-related research; HLA class II-associated disease; transplantation matching; recombination hot spot identification; linkage disequilibrium study; human; microsatellite.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human microsatellite
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                                                                                                       Sequence
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                                                   18;
                                                                Similarity
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                                                   Conservative
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                                                                                                      A; 6 C; 5
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90.0%;
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                                                                                                       G; 3 T; 0
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0; Mismatches
                                                                Score 16.8;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer
                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    #2
                                                                                                         0 Other
                                                                  .6e+03;
                                                                             DB 1;
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06-NOV-2003

(first entry)

Homo

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RESULT 1357
ADB81564/c
ID ADB8156
XX
AC ADB8156
XX
DT 04-DEC-
XX
DE Antiser
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                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to a novel antisense compounds that modulate the capression of oestrogen receptor beta (ERSB). Oestrogen is a steroid commone that exerts a wide range of effects throughout the human body being primarily involved in female sexual maturation. Additionally, component that exerts a wide reproductive tissues, is known to be convery oestrogen targets male reproductive tissues, is known to be converted to bone maintenance and plays a protective role in the coardiovascular system. This hormone receptor, ERSB (also known as ER coardiovascular system. This hormone receptor, ERSB (also known as ER coardiovascular system. This hormone receptor, ERSB (also known as ER coardiovascular system. This maturation cells indicating an involvement in coasse, uterine leiomyomata and neoplasms of the kidney. Furthermore, coardingly, the selective inhibition of ERSB by coardingly, the selective inhibition of ERSB by coardingly, the selective inhibition of ERSB by coardingly captured to the sinvention could provide a therapeutic target for the treatment of cancer, as well as other ERSB-coardingly selective inhibition of ERSB by coardingly selective inhibition of ERSB by coardingly selective sequence is the chimeric human captured to antisense oligo used to inhibit expression of human ERSB, the aim of the cinvention. Note that it has two terminal five nucleotide 2-methoxyethyl coardingly separated by a ten deoxynucleotide gap. The coligonucleotide backbone is phosphorothioate throughout
                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oestrogen receptor beta; ERSB; steroid hormone; female sexual maturation; bone maintenance; cardiovascular system; ER beta; oestrogen receptor 2; ERS2; Alzheimer's, uterine leiomyomata; cytoostatic; kidney neoplasm; ss; cellular proliferation; cancer; human; antisense; chimeric.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides for modulating estrogen receptor beta gene expression, particularly useful for treating cancers, specifically leiomyoma, pancreatic cancer, prostate cancer, breast cancer, bone cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chimeric antisense oligonucleotide ISIS 192360 to inhibit human ESRB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dobie KW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-DEC-2001; 2001US-00005058.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-DEC-2002; 2002WO-US039200.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003050133-A1.
  Antisense oligo (SeqID 81) used to inhibit human EIF2C1 DNA.
                                         04-DEC-2003
                                                                                                                      ADB81564 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 3; Page 81; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                            730
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                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                        GTAGCTGGGATTACAGGTGC 1
                                                                                                                                                                                                                                                        GTAGCTGGGACTACAGGCGC 749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Roach MP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                 BP; 5
                                                                                                                                                                                                                                                                                                      Conservative
                                       (first entry)
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                                                                                                                      DNA;
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                                                                                                                                                                                                                                                                                                                                                                                 G; 4 T;
                                                                                                                                                                                                                                                                                                                    Score 16.8; DB 1;
Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                   U; 0 Other;
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04-DEC-2003 ADB81567;

(first entry)

ADB81567 standard; DNA; 20

BP.

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RESULT 1358
ADB81567/c
ID ADB8156
XX
AC ADB8156
XX
AC ADB8156
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O4-DEC-
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the expression of human eukaryotic translation initiation factor 2C 1 (EIF2C1). EIF2C1 is located on chromosome 1934-35, and is also known as Co-eIF2C, eIF2C, Golgi ER protein 95kDa, GERp95 and Q99. It is an intracellular membrane associated protein thought to be involved in cellular differentiation, such that altered expression of EIF2C1 can affect cell growth, morphology and tumourigenicity. Accordingly, antisense oligonucleotides that inhibit the expression of EIF2C1 in cells or tissues can be used in gene therapy to treat various conditions including hyperproliferative disorders, familial hypercholesterolaemia and cancer, as well as polycystic kidney disease, cystic fibrosis and progeriod syndrome. As such, the oligos of the present invention can be described as having cytostatic and antisense oligo used to inhibit expression of the human eukaryotic translation initiation factor 2C 1 (EIF2C1) DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New compound, having a sequence targeted to a nucleic acid encoding human collapsin response mediator protein 2, useful for preparing a composition for treating hypercholesterolemia or hyperproliferative disorder, e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 3; Page 77; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-449448/42.
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                                                                                                                                                                                                                                                                                                                             640 TCACCCAGGCTGGAGTGCAG 659
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human eukaryotic translation invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 4 A; 8 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       relates to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag=
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                                                                                                                                                                                                                                                                                                                                                                                        0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 16.8;
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Ward

; pituitary; pituitary disease; joint disease; rheumatoid arthritis;

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RESULT 1359
ADB90595
ID ADB9059
XX
AC ADB9059
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                                                                                                                                                                                                               Query Match
Best Local (
                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense; ss; human; eukaryotic translation initiation factor 2C : EIF2C1; CO-EIF2C; EIF2C2; GO1gi ER protein 95kDa; GERp95; Q99; gene therapy; hyperproliferative disorder; familial hypercholesterolaemia; cancer; polycystic kidney disease; cystic fibrosis; progeriod syndrome; cytostatic; antilipaemic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New compound, having a sequence targeted to a nucleic acid encoding human collapsin response mediator protein 2, useful for preparing a composition for treating hypercholesterolemia or hyperproliferative disorder, e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Page 77; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-449448/42
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                                                                                                                                                                                                                                                                   Sequence
                                        ADB90595
                                                                                                                                                             661
                                                                                                                           20
                                                                                                                                                                                                                                                                                                      human eukaryotic invention.
                                                                                                                                                                                                               Similarity
                                                                                                                                                       GGCGCAATCTTGGCTCACTG 680
                                      standard;
                                                                                                                           GGCACGATCTTGGCTCACTG 1
                                                                                                                                                                                                                                                                   20 BP; 5
                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod base= OTHER /mod="0THER" phosphorothioate backbone, where 1-5 and /note= "OTHER= phosphorothioate backbone, where 1-5 and 16-20 are 2' methoxyethyl nucleotides. All cytidines are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SeqID 84) used to inhibit human EIF2C1 DNA
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                                        DNA;
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                                          BP.
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                                                                                                                                                                                                                                                                      G; 3 T;
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Pred. No. 1.
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                                                                                                                                                                                                                                                                      Other;
                                                                                                                                                                                                                                 Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1360
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention comprises the amino acid and coding sequences of human proteins (PGSF1a, PGSF1b, PGSF2 and Pi-a) that are specifically expressed in the human pituitary. The DNA and protein sequences of the invention are useful for the diagnosis and treatment of pituitary disease (e.g. autoimmune pituitary inflammation) and joint diseases (e.g. rheumatoid arthritis). The present DNA sequence was used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-SEP-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-DEC-2003 (first entry)
                                                                                                                                                                                                Cytostatic; antisense therapy; co-repressor; RE1 silencing transcription factor; COREST; antisense oligonucleotide; developmental; hyperproliferative; disorder; neuronal cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
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                        modified_base
                                                                                                           modified_base
                                                                                                                                                                                                                                                                          Human CoREST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Proteins expressed specifically in human pituitary and antibodies to th for diagnosis and treatment of pituitary-associated and joint diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tatsumi K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NISC-)
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                                                                                                                                                                                                                                                                                                           01-JAN-2004
                                                                                                                                                                                                                                                                                                                                              ADC89590
                                                                                                                                                                                                                                                                                                                                                                                  ADC89590 standard; DNA;
                                                                                                                                                                sapiens
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18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GCCTCAGCCTCCCGAGTATC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fla; PGSF1b; PGSF2; Pi-a;
pituitary inflammation; *
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                        antisense oligonucleotide #ISIS 165030
                                                                                                                                                                                                                                                                                                           (first entry)
                                      /note= "phosphorothioate backbone"
/note= "all cytidines are 5-methyl
                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7; 58pp; Japanese.
                                                                                          /*tag=
                                                                             mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;
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Pred. No. 1
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Mismatches

.6e+03;

Length Indels

20;

2

0,

Gaps

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0 Other; DB 1;

Okubo K;

them

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are 5-methylcytidines"

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ADC89591/c
ID ADC8959
XX ADC8959
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XX develop
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FT modifie
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a new antisense compound comprising 8-50 nucleobases in length targeted to a nucleic acid molecule encoding a corepressor for REI silencing transcription factor (COREST), where the compound specifically hybridises with and inhibits the expression of COREST. The COREST antisense oligonucleotide has any of 72 specifically claimed sequences of 20 bp, given in the specification. The methods and compositions of the present invention are useful for the diagnosis, prevention and/or treatment of diseases or conditions associated with aberrant expression or activity of COREST, such as a developmental disorder and/or a hyperproliferative condition like neuronal cancer. The current sequence represents an antisense oligonucleotide for the inhibition of human COREST MENNA levels. Nucleotides of the invention have
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense oligonucleotide compounds, useful for the diagnosis, prevention and/or treatment of conditions with aberrant expression or activity of COREST, such as developmental and/or hyperproliferative
                                                                                                                                                       Human CoREST
                                                                                                                                                                                      01-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-256431/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-AUG-2001; 2001US-00920671
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-JUL-2002; 2002WO-US024370
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                         modified_base
                                                                                                 developmental;
                                                                                                                             Cytostatic;
                                                                                                                                                                                                                                             ADC89591 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC.
                                                                                                              silencing
                                                                     sapiens
                                                                                                                                                                                                                                                                                                                                            383
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                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                               20
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                                                                                                                                                                                                                                                                                                                  CCTCCCAAAGTGCCAGGATT 20
                                                                                                                                                                                                                                                                                                                                                                                                                             B₽;
                                                                                               antisense therapy; co-repressor; og transcription factor; COREST; al; hyperproliferative; disorder;
                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                       antisense oligonucleotide #ISIS 165031.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Freier
                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                             v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "
 /mod
                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          note= "2'-0-methoxyethyls (2'MOE) wing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                             ð
                           . 20
                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          81; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SM;
                                                                                                                                                                                                                                                                                                                                                                                 1.7%;
90.0%;
_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                               7
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                                                                                                                                                                                                                                                                                                                                                                                                                             C; 4 G; 4 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "2'-O-methoxyethyls (2'MOE) wing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       a
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                                                                                                                                                                                                                                                                                                                                                                                   Score 16.8; DB 1;
Pred. No. 1.6e+03;
                                                                                                            co-repressor;
ctor; COREST; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                             Ġ.
                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other;
                                                                                                 neuronal
                                                                                                                                                                                                                                                                                                                                                                                                 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                 cancer;
                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                       Gaps
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RESULT 1362
ADD21697/c
ID ADD2169
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                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a new antisense compound comprising 8-50 nucleobases in length targeted to a nucleic acid molecule encoding a corepressor for REI silencing transcription factor (COREST), where the compound specifically hybridises with and inhibits the expression of COREST. The COREST antisense oligonucleotide has any of 72 specifically claimed sequences of 20 bp, given in the specification. The methods and compositions of the present invention are useful for the diagnosis, prevention and/or treatment of diseases or conditions associated with aberrant expression or activity of COREST, such as a developmental disorder and/or a hyperproliferative condition like neuronal cancer. The current sequence represents an antisense oligonucleotide for the inhibition of human COREST mRNA levels. Nucleotides of the invention have
                       antisense oligonucleotide; human; mdm2; hyperproliferation; hyperproliferative disorder; cancer; psoriasis; fibrosis; atherosclerosis; restenosis; apoptosis modulation; p21; ss; 2'-methoxyethoxy-residue; phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense oligonucleotide compounds, useful for the diagnosis, prevention and/or treatment of conditions with aberrant expression or activity of COREST, such as developmental and/or hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                  Human mdm2 antisense oligonucleotide #260.
                                                                                                                                                                                          ADD21697 standard;
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-JUL-2002; 2002WO-US024370.
                                                                                                                                                             ADD21697;
                                                                                                                                                                                                                                                                                                                                                                                                                  2-MOE wings and a deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-FEB-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS
                                                                                                                                L5-JAN-2004
                                                                                                                                                                                                                                                                                              722 CCTCCTGAGTAGCTGGGACT 741
                                                                                                                                                                                                                                                                   20 CCTCCCGAGTAGCTGGGATT 1
                                                                                                                                                                                                                                                                                                                            18;
                                                                                                                                                                                                                                                                                                                                                                                     BP; 5 A; 6 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001US-00920671.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ID NO 82; 145pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Freier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PHARM INC.
                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "phosphorothioate backbone"
/note= "all cytidines are 5-methyl-
1. .5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mod
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note=
                                                                                                                                                                                            DNA;
                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         _base= OTHER
e= "2'-O-methoxyethyls
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  base= OTHER
e= "2'-O-methoxyethyls
                                                                                                                                                                                                                                                                                                                                          90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                  gap.
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                                                                                                                                                                                                                                                                                                                                          Score 16.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                           .6e+03
                                                                                                                                                                                                                                                                                                                                                       DB 1; .Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (2'MOE) wing"
                                                                                                                                                                                                                                                                                                                               Indels
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Homo sapiens

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ADD21700/c
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the human mdm2 gene. The antisense oligonuclectides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonuclectides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonuclectides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonuclectide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel antisense compound targeted to 5' untranslated region, coding region, or intron:exon junction of nucleic acid molecule encoding mdm2, useful for treating e.g. cancer, psoriasis or restenosis by inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Manoharan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Miraglia LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-DEC-2002; 2002WO-US038281.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2003048315-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 6 A; 4 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 9; SEQ ID NO 262; 289pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-JUN-2003
                                                                                                                                                                          hyperproliferative disorder; cancer; psoriasis; fibrosis; atherosclerosis; restenosis; apoptosis modulation; p21; s 2'-methoxyethoxy-residue; phosphorothioate backbone.
                                               04-DEC-2001; 2001US-00005344
                                                                         02-DEC-2002; 2002WO-US038281
                                                                                                                           WO2003048315-A2
                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                  antisense oligonucleotide; human; mdm2; hyperproliferation;
                                                                                                                                                                                                                                           Human mdm2 antisense oligonucleotide #263.
                                                                                                                                                                                                                                                                     15-JAN-2004
                                                                                                                                                                                                                                                                                                                         ADD21700 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present sequence contains 2'-methoxyethoxy-residues and has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention comprises antisense oligonucleotides which are targeted to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2003-577263/54
                                                                                                                                                                                                                                                                                                                                                                                                                  213
                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                                                                                                                            18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                              GGTCTCGAACTCCCGACCTC 232
                                                                                                                                                                                                                                                                                                                                                                                         GGTCTCGATCTCCTGACCTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2001US-00005344.
                                                                                                                                                                                                                                                                     (first entry)
                         PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nero PS,
Nero PS,
                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Graham MJ,
                                                                                                                                                                                                                                                                                                                          20
Graham MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 16.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Monia
Monia
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₽₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Koller E,
Koller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chiang
Chiang
                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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RESULT 1364
ADD21693/c
ID ADD2169
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Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the human mdm2 gene. The antisense oligonucleotides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonucleotides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonucleotides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel antisense compound targeted to 5' untranslated region, coding region, or intron:exon junction of nucleic acid molecule encoding mdm2 useful for treating e.g. cancer, psoriasis or restenosis by inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 9; SEQ ID NO 265; 289pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mdm2 expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-577263/54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Manoharan M;
                                                                Novel antisense compound targeted to 5' untranslated region, region, or intron:exon junction of nucleic acid molecule encuseful for treating e.g. cancer, psoriasis or restenosis by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 2 A; 3 C; 11 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence contains 2'-methoxyethoxy-residues phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention comprises antisense oligonucleotides which are targeted
                                                                                                                                                        Miraglia LJ,
Manoharan M;
                                                                                                                                                                                                                                                                                                                                                                         2'-methoxyethoxy-residue;
                                                                                                                                                                                                                                                                                                                                                                                      hyperproliferative disorder; cancer; psoriasis; fil atherosclerosis; restenosis; apoptosis modulation;
                                                                                                                                                                                                                                                                                                                                                                                                                     antisense oligonucleotide; human; mdm2; hyperproliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADD21693 standard; DNA; 20
                                                                                                                                                                                                                             04-DEC-2001;
                                                                                                                                                                                                                                                         02-DEC-2002; 2002WO-US038281
                                                                                                                                                                                                                                                                                                                 WO2003048315-A2
                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JAN-2004
                        Claim 4; SEQ ID NO 258; 289pp; English
                                                      mdm2 expression
                                                                                                                            WPI; 2003-577263/54
                                                                                                                                                                                                  (ISIS-) ISIS
                                                                                                                                                                                                                                                                                      12-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         842 GCCTGCCTCGGCCTCCCAAA 861
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18;
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                                                                                                                                                                                                  PHARM INC.
                                                                                                                                                                                                                              2001US-00005344.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                       Nero PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 oligonucleotide #256
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                                                                                                                                                                       Graham MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ΒP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 16.8; DB 1;
Pred. No. 1.6e+03;
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                                                                                                                                                                        Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                        Koller E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
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                                                                      encoding mdm2
by inhibiting
                                                                                                                                                                          Chiang MY,
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The invention comprises antisense oligonucleotides which are targeted to

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ADD21686/c

XX

XX

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ADD21

XX

DT 15-J

XX

DT 15-J

XX

DT 15-J

XX

DT Huma

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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                               the human mdm2 gene. The antisense oligonucleotides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonucleotides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonucleotides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonucleotide of the invention. The present sequence contains 2'-methoxy-residues and has a
                                                                                                                                                                                                                                                                                                                                                           Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel antisense compound targeted to 5' untranslated region, coding region, or intron:exon junction of nucleic acid molecule encoding mdm2 useful for treating e:g. cancer, psoriasis or restenosis by inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-577263/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Miraglia LJ,
Manoharan M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-DEC-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hyperproliferative disorder; cancer; psoriasis; fik atherosclerosis; restenosis; apoptosis modulation;
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phosphorothioate backbone.
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                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 251; 289pp; English.
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                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                cytostatic, neuroprotective, mostropic, antiarteriosclerotic, antiarthritic and nephrotropic activities and may be useful in identifying compounds that inhibit the induction of genes involved in viral infection, cancer, renal diseases or age-related diseases including Alzheimer's disease, atherosclerosis or arthritis, such genes being induced by cyclin-dependent kinase inhibitors. Furthermore, The constructions have gene therapy applications. The current sequence is that of the PCR primer which was used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New recombinant expression construct for identifying and modulating expression of genes regulated by cyclin-dependent kinase inhibitors, as genes involved in viral infection, cancer, renal diseases or age-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                virucide; cytostatic; neuroprotective; nootropic; antiarteriosclerotic;
antiarthritic; nephrotropic; viral infection; cancer; renal;
age-related disease; Alzheimer's; atherosclerosis; arthritis;
                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        encoding a reporter gene operably linked to a promoter from a man
viral or cellular gene induced by a cyclin dependent kinase (CDK)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inhibitor. The construct of the invention demonstrates virucide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel recombinant expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-AUG-2001; 2001US-0315791P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  recombinant expression construct; cyclin-dependent kinase inhibitor;
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                                                   723 CTCCTGAGTAGCTGGGACTA 742
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human RANTES oligonucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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                                                                                                                                Sequence
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                                                                                   Similarity
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ABZ92735
                                                                                                                                                                                                                                                                                                                                                                 consent airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention thas antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a cimmunosuppressive, and cytostatic activity. The composition may have a cimmunosuppressive, and cytostatic activity. The composition may have a cimmunosuppressive, and cytostatic activity. The composition may have a consisting a respiratory, lung or malignant disease or condition, also consider the prophylactic or therapeutic respiratory effect of an continflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodollation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, constitution, inflammation, lung allergies, or a respiratory disease or condition. The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel pharmaceutical composition, which has if the first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5 or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or corresponding control of steroid or corresponding control or correspondin
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Tang L,
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                                                                                                                                                     Sequence 20
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bt active agent comprising an oligonucleotide antisense to the
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L, Shahabuddin
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Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a
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CC lung inflammation, lung allergies, or a respiratory disease or condition.

CC Note: The sequence data for this patent is not represented in the printed as pecification, but was obtained in electronic format directly from WIPO as formation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or corresponding controls.
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                                                                 Sequence 20
                                                                                                                       lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the print specification, but was obtained in electronic format directly from WIPC specification, but was obtained in electronic format directly from WIPC.
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                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                            lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                            BP; 19 A; 0 C; 0 G; 1 T; 0 U; 0 Other;
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                                                                                                                      Conservative
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L, Shahabuddin
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                                                                                                                                                                               Score 16.8;
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                                                                                                         Matches
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                                                                                                                                                                                                                                                            lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                  Sequence 20 BP; 19 A; 0 C; 0 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel pharmaceutical composition,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ubiquinone
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18; Conserv
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, Tang L,
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                                                                                                         Conservative
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L, Shahabuddin S;
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                                                                                                                                  90.0%;
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Matches 18
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                                                                                                                                                                                                                                                                                                                                                                    lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 5105; 872pp; English
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                                                                                                                                                                                                                                                                                                    Sequence
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, Shahabuddin
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                                                                                                                                                                                                                                DB 1;
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5' and 3' intron-exem junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of beginning or the composition of the prophylactic continuous control of the subject.
                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5, or 3, end genomic flanking regions, or regions within 2-10 nucleotides of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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  Sequence 20
                                                           lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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    BP; 5 A; 2 C; 8 G; 5 T; 0 U; 0 Other;
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L, Shahabuddin
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Query Match Best Local Matches 1

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Conservative

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Mismatches

Similarity

90.0%;

Score 16.8; Pred. No. 1

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Tang L,
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L, Shahabuddin
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Miller S,
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                                                                                                                                               Sequence 20 BP; 3 A; 5 C; 6 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                 lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Some sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel pharmaceutical composition, which I first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking region, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides,
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                                                                                                       lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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CC Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO or at ferm when in the printed in the printed of the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 14305; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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                                                                                                                                                                                                                                                                                                                                                                                                                             CC junctions of genes encoding a polypeptide associated with lung and/or cc antiinflammatory steroid and ubiquinone. A composition of the invention cc antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a crown antisense gene therapy. The composition is useful for treating or cc use in antisense gene therapy. The composition is useful for treating or cc reventing a respiratory, lung or malignant disease or condition, also composition a subject, for reducing or depleting levels of an antiinflammatory steroid in a subject, for reducing or depleting levels of adenosine cf. or reducing sensitivity to adenosine, reducing levels of adenosine conformation, lung surfactant in a subject's tissue, or treating bronchoconstriction, cc lung surfactant in a subject's tissue, or respiratory disease or condition. Cc lung inflammation, lung allergies, or a respiratory disease or condition. Cc lung inflammation, but was obtained in electronic format directly from WIPO case for the printed and the printed control of the printed and the printed and the printed are the printed and the printed and the printed are the printed and the printed and the printed are the printed and the printed and the printed are the printed and the printed are the printed and the printed are the printed and the printed and the printed and the printed and the printed are the printed and the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ÄBZ99109;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; antisense; lung dysfunction; nasal airway dysfunction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PDE4C oligonucleotide sequence.
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Tang L,
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L, Shahabuddin
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                                                                                                                                                                                                                                                                   immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or
                                                                                                                                                                                                                                                                                                                                                                                                junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonucleotide antisense to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure;
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antisense gene therapy; respiratory; lung; adenosine sensitivity;
adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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                                                                      18;
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Tang L,
                                                                                                                                            BP; 4 A; 10 C; 2 G; 4 T; 0 U; 0 Other;
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                                                                      Conservative
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L, Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, inmunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or
                                                                                                                                                                                                                                                                                                 lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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                                                          GGCTGGAGTGCAGTGGCGCA 666
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                                                                                                                                                                                                                                                          lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                          l Similarity
18; Conser
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Tang
  GTATGTTTAGTAGAGACGGG 20
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CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC antiinflammatory steroid and ubiquinone. A composition of the invention CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an antisnflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine creceptor, producing bronchodulation, increasing levels of adenosine surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. CC Note: The sequence data for this patent is not represented in the printed constriction, but was obtained in electronic format directly from WIPO at the proposition of the printed constriction in the printed c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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  Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 13147; 872pp; English
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Tang L,
        BP; 2 A; 6 C; 8 G; 4 T; 0 U;
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L, Shahabuddin
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           0 Other;
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Matches Query Match Best Local

1 Similarity
18; Conserv

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                                                                                                                                                     Sequence 20
                                                                                                                                                                                                         lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure;
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                                                                                            Local Similarity
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                                  1033 GCTGGGATTACGGGCACCTG 1052
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Tang L,
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                                                                          Conservative
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Miller S,
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                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammation; respiratory disease;
                                  195
                                                                      18;
                                                                                       Similarity
                                                                                                                                          20
 CGCCATGTTGGCCAGGCTGG
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Tang
                                                                                                                                          BP;
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                                                                      Conservative
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Shahabuddin
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                                                                                                                                          6 C;
                                                                                   1.7%;
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                                                                                                                                          G; 4 T;
                                                                    <u>,,</u>
                                                                                     Score 16.8;
Pred. No. 1
 20
                                  214
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S;
                                                                      Mismatches
                                                                                                                                          0 U; 0 Other;
                                                                                       1.6e+03;
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                                                                                                          BB
                                                                                                      Length 20;
                                                                        Indels
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ABZ92730

ABZ92XX,
AC, ABZ92XX,
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Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic;
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                                                                                                                                                                                                                  lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                       Sequence 20
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Similarity
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Tang L,
                                                                                                           BP; 2 A; 4 C; 8
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L, Shahabuddin
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90.0%;
                                                                                                           G; 6 T; 0 U; 0 Other;
                                   Score 16.8;
Pred.
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   No.
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                                   Length
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Matches

18;

Conservative

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Indels

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Gaps

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201 GTTGGTCAGGCTGGTCTCGA 220

GTTGGCCAGGCTGGTCTTGA 20

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                                                                                                         Matches
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                                                                                                                                                                                                                                                                                             lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Мусе JW,
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                                                                                                                                                                                                                                                                ftp.wipo.int/pub/published_pct_sequences
                                                   646
                                                                                                         18;
                                                                                                                                      Similarity
                                  AGGCTGGAGTGCAGTGGCGC 665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Li Y, Sa
Tang L,
                                                                                                                                                                                                                   BP; 4 A; 3 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                         Conservative
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L, Shahabuddin
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                                                                                                                                   90.0%;
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                                                                                                                                      Pred. No.
                                                                                                                                                          Score 16.8;
                                                                                                         Mismatches
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                                                                                                                                      1.6e+03
                                                                                                                                                             DB 1; Length 20;
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                                                                               Matches
                                                                                                    Best Local
                                                                                                                     Query Match
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Miller S,
                                                                                                                                                                                            lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                             Sequence 20 BP; 5 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; antisense; lung
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-APR-2002; 2002WO-US013135
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                                         472 AGGATGAAGTGCAGTGGTGT 491
20
                                                                               18;
                                                                                                    Similarity
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, Tang L,
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L, Shahabuddin S;
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                                                                                                    90.0%;
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                                                                                                    Pred. No.
                                                                                                                       Score 16.8; DB 1; Length 20;
                                                                                 Mismatches
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                                                                                                    1.6e+03
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RESULT 1394

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ABZ99061

ABZ9
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                                                                                                                                               Query Match
Best Local :
                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                        lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                     Sequence
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Tang L,
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AGTGATTCTCCTGCCTCAGC
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The invention relates to a novel pharmaceutical composition, which has a cfirst active agent comprising an oligonucleotide antisense to the cinitiation codon, coding region, 5' or 3' end genomic flanking regions, coding region, so regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or compassion and a second active agent comprising an compassion and a second active agent comprising an committed antinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a communosuppressive, and cytostatic activity. The composition may have a compassion and cytostatic activity. The composition, also compressive, and cytostatic activity. The composition, also composition are preparatory, lung or malignant disease or condition, also composition are respiratory at the prophylactic or therapeutic respiratory effect of an composition and the prophylactic or therapeutic respiratory effect of an initial mmatory steroid in a subject, for reducing or depleting levels of conformation, lung surfactant in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of biquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, continuing antipal and the printed specification, but was obtained in electronic format directly from WIPO at ferm wino intrombinal had not agenciance.
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                                                                                                           Sequence 20
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   936 TCTGTTACCCAGGCTGGAGT 955
                                       18;
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Tang L,
                                                                                                             BP; 2 A; 4 C; 8 G; 6 T; 0 U;
                                       Conservative
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                                                                                                                                                                                                                                                                      Sequence 20
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                                                                  CTCAGCCTCCCAAGTAGCTG 561
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                                                                                                                                     Conservative
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cc first active agent comprising an oligonucleotide antisense to the cc initiation codon, coding region, 5 or 3 end genomic flanking regions, cc 5 and 3 intron-exon junctions, or regions within 2-10 nucleotides of cc junctions of genes encoding a polypeptide associated with lung and/or cc masal airway dysfunction and a second active agent comprising an cc antiinflammatory steroid and ubiquinone. A composition of the invention cc antiinflammatory, antiallergic, antiasthmatic, hypotensive, cc immunosuppressive, and cytostatic activity. The composition may have a cc use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also cf or enhancing the prophylactic or therapeutic regiratory effect of an cc antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine cc receptor, producing bronchodilation, increasing levels of twing inflammation, lung allergies, or a respiratory disease or condition. CC ung inflammation, lung allergies, or a respiratory disease or condition. CC specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Miller S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antisense; lung dysfunction; nasal airway dysfunction;
                                                                      18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EPIGENESIS PHARM INC
                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Li Y, Sa
, Tang L,
                                                                                                                                          20 BP; 4 A; 7 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 5095; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sandrasagra A, Katz E, L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   respiratory disease; ds
                                                                                      90.0%;
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                                                                      0;
                                                                                        Pred.
                                                                                                         Score 16.8; DB 1; Length 20;
                                                                       Mismatches
                                                                                        No.
                                                                                        1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pabalan J,
                                                                       ب
                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Aguilar
                                                                      <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     which has a
                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         therapy;
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ABZ89860/c

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IID ABZ898

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Human

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Human

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                                                                                                                                                                                                                                                        tirst active agent comprising an oligonucleotide antisense to the contribution codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC enthancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing bensitivity to adenosine, reducing levels of adenosine creceptor, producing bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition. CC Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at fig. wipo.int/pub/published_pot_sequences
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                                                                                                              Query Match
Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABZ89860 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human oligonucleotide sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID NO 5102; 872pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-229219/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-APR-2001; 2001US-0286137P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-APR-2002; 2002WO-US013135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200285308-A2
                                                                                                                                                                                                                               Sequence
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Tang
                                                       CACTGCAACCTCTGCCTCCC 695
   CACTGCAACCTCCGCCTTCC
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                                                                                                                                                                                                                               BP; 4
                                                                                                              Conservative
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L, Shahabuddin
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                                                                                                                                             1.7%;
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C;
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                                                                                                                                             Score 16.8;
Pred. No. 1
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                                                                                                                                                                                                                                  3 T; 0 U;
                                                                                                                    Mismatches
                                                                                                                                                1.6e+03
                                                                                                                                                                                                                                  0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pabalan J,
                                                                                                                                                                              DB 1;
                                                                                                                                                                        Length
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RESULT 1400

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ACA88890/c
ID ACA888
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AC ACA888
AC ACA888
DT 08-JUL
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HOMO 8
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          genetic markers according to a heterozygosity index of 0.5 or greater. Selecting and amplification of genetic markers are useful as targets for nucleic acid sequence amplification, for genetic testing or facilitating multiplex PCR amplification from limiting amounts of target nucleic acid. The methods are also useful for improving genetic diagnostic and screening methods, such as prenatal diagnostic testing, foetal sex determination or genetic identification, e.g. DNA profiling or DNA fingerprinting. The nucleic acid sequence amplification is also useful in forensic analysis of degraded, old, ancient and difficult tamples that are difficult to amplify and identify. This sequence represents a PCR primer used in the selection and amplification of genetic markers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Selecting genetic markers as targets for nucleic acid sequence amplification, useful for improving genetic testing, e.g. fetal sex determination, comprises selecting each of the genetic markers according
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genetic marker selection; multiplex PCR amplification; prenatal diagnostic testing; foetal sex determination; genetic identification; DNA profiling; DNA fingerprint; forensic analysis; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Selection and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-JUL-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention describes a method of selecting genetic markers as targets for nucleic acid sequence amplification comprising selecting each of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 36; Page 39; 64pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-OCT-2001; 2001AU-00008234.
12-OCT-2001; 2001AU-00008235.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-OCT-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-APR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
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                               Human Y chromosome non-recombining region polymorphic fragment
                                                                                      03-JUN-2004
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                                                                                                                                            ADM65766;
                                                                                                                                                                                                   ADM65766 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                       CAGCCTCCCAAAGTGCCTGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matthews
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G; 4 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 16.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.6e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cc male comprising obtaining a nucleic acid sample from the male, and comprising obtaining a nucleic acid sample from the male, and condition of the ethnic origin of the male, using at least two polymorphic markers in the nucleic acid sample condition from the primer pairs given in the specification. Also described is called the ethnic origin of an individual; determining the ethnic origin of an individual; determining the ethnic contiguous bases including at corigin of a human male individual; an isolated nucleic acid segment of a contiguous bases including at called the polymorphic sites given in the specification; nucleic conditions of the polymorphic sites given in the specification; and determining the paternity of a human male condition of the polymorphic sites given in the specification; and determining the ethnic origin of a conditional. The method is useful for determining the ethnic origin of a condition of the specification; and determining the ethnic origin of a condition of the number of the non-recombining region of the human conditions of the human conditions (and the primer pairs of the number of the number of conditions) continued the condition of the human conditions (and the primer pairs of the number of the number of the number of the number of the human conditions).
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Matches 18
              Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hyportension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Determining the ethnic origin of a male by obtaining a nucleic acid sample from the male and identifying at least two polymorphic market the nucleic acid sample indicative of the ethnic origin of the male
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Y chromosome; paternity non-recombining region;
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                                                                                                                                                   Human PDE4C-derived oligonucleotide SEQ ID 14350.
                                                                                                                                                                                                                                                                         ABD32139 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention describes a method of determining the ethnic origin of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 24; Page 66; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-843259/78
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emphysema; chronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   determine ethnic origin of a male.
                                                                                                                                                                                              29-JUL-2004
                                                                                                                                                                                                                                    ABD32139;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               origin determination; polymorphic site determination;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OEFNER P J.
UNDERHILL P A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                     TGTTGGTCAGGCTGGTCTCG 219
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                                                                                                                                                                                          (first entry)
obstructive pulmonary disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7 C; 3 G; 1 T; 0 U; 0 Other;
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human; NRY; polymorphic fragment;
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Pred.
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No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20;
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RESULT 1403 ABD32140 ID ABD3214 XX AC ABD3214

ABD32140

standard; DNA; 20

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expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. Combetion of the containers of the corresponding mRNA. Combetion also describes a kit, that comprises: (a) a delivery combetion also describes a kit, that comprises: (a) a delivery combetion of the invention activity and for use of the kit. The composition combetion has antiallergic, antiinflammatory, antiasthmatic, combetion is useful for preventing or combetion comprises oligo and is administered to reduce the production composition comprises oligo and is administered to reduce the production correduce the amount of target polypeptide present in the lungs. The composition composition, and/or bronchoconstriction and/or lung confilmmation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vestication.
                                                                                                                        Matches
                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                 inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, aller reducing adenosine sensitivity, levels of adenosine (A) or
                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      surfactant depletion or hyposecretion, when administered to a mammal. oligonucleotides are derived from a gene encoding or regulating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-APR-2001; 2001US-0286036P
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                                                                                                                     Local Similarity
les 18; Conserv
                                                       535 CTCCTGCCTCAGCCTCCCAA 554
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, Tang L,
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CTCCCACCTCAGCCTCCCAA 20
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L, Shahabuddin
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                                                                                                                                                                                                                                                                                                                 effects due
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Pred. No. 1
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                                                                                                                                                  1.6e+03
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                                                                                                                        Gaps
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Query Match
Best Local Similarity
  1.7%;
Score 16.8;
Pred. No. 1.
   1.6e+03;
            DB 1;
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beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; surfactant depletion; immunosuppressive; cytostatic; cystic fibrosis; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; Human PDE4C-derived oligonucleotide 29-JUL-2004 (first entry) SEQ ID 14351

WO200285309-A2

31-OCT-2002

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

ŝ Li Y, Tang ŗ Sandrasagra A, L, Shahabuddin Katz S; Ħ Pabalan J, Aguilar Ď

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and

ij NO 14351; 763pp; English.

Comprising oligonuclectides, effective for alleviating comprising oligonuclectides, effective for alleviating comprising oligonuclectides, effective for alleviating allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, considered to a mammal of the coligonuclectides are derived from a gene encoding or regulating considered to a mammal. The coligonuclectides are derived from a gene encoding or regulating considered to a mammal of the coligonuclectides are derived from a gene encoding or regulating considered with lung airway or lung considered to the corresponding mRNA. Considered to responding mRNA. Considered to reduce, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, and beta-adrenergic agonist. The composition is useful for preventing or considered composition comprises oligo and is administered to reduce the production consistion comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition construction, and/or bronchoconstriction and/or lung construction, and/or bronchoconstriction and/or lung consisting a respiratory. Antiatory consisting to the discress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation, respiratory construction, allergies, asthma, impeded respiration, respiratory construction and asthmatic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonuclectides into products that free adenosine into the system considered considered the system of the system of the system of the system of the system o

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0 Other;

Matches 381 AGCCTCCCAAAGTGCTGGGA 400 18; AGCCTCCCAAAGTACCGGGA Conservative 0 20 Mismatches *ې* Indels 0 Gaps

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ABD32094 standard; DNA; 20 ΒP

Human PDE4C-derived oligonucleotide SEQ ID 14305.

29-JUL-2004

(first entry)

respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; cystic fibrosis; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hyportension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;

Homo sapiens.

WO200285309-A2

31-OCT-2002

23-APR-2002; 2002WO-US013143

2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

Miller **Мусе JW** ŝ Li Y, Tang ŗ Sandrasagra A, 5, Shahabuddin Katz E, Pabalan 'n Aguilar Ö

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent

Claim 15; IJ NO 14305; 763pp; English

CC surfactant depletion or hyposecretion, when administered to a mammal. The capture of dysfunction or a target polypeptide associated with lung airway or lung construction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, considering a respiratory, lung or malignant disease. The administered composition composition composition composition is useful for preventing or constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, considered composition comprises oligo and is useful for preventing or composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, allergies and/or surfactant hypoproduction are associated inflammation. allergies and immeded respiration. bronchoconstriction, respiratory tract inflammation, allerg reducing adenosine sensitivity, levels of adenosine (A) or comprising oligonucleotides, effective for alleviating This invention describes a novel composition inflammation, allergies, asthma, impeded respiration, krome, pain, cystic fibrosis, allergic rh (a) a first active agent, allergies and (A) receptors, pulmonary

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ABD32118
ID ABD3211
AC ABD321
AC ABD
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Best Local Similarity
Matches 18; Conserv
This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human PDE4C-derived oligonucleotide SEQ ID 14329.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              bronchodilating agent.
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Tang
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L, Shahabuddin
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Matches 18
                                                                                                                                                                                                                                                                                    Nyce JW,
Miller S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the oligonucleotides into products that e.g., lung, brain, heart, kidney, etc, t prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                             24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-APR-2002; 2002WO-US013143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-OCT-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           rejection; ss; primer.
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Pred. No. 1
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Claim 15; SEQ ID NO 7967; 763pp; English

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targenucleic acids associated with lung airway or lung dysfunction

dysfunction,

targeted to

WPI; 2003-093058/08

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CC The invention also describes a kit, that comprises: (a) a delivery construction also describes a kit, that comprises: (a) a delivery consistency in separate containers, (b) the oligonucleotidee, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, of the invention has antiallergic, antiinflammatory, antiasthmatic, and consistency, lung or malignant disease. The administered or composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to cor availability, or to increase the degradation of the target mRNA or to construction, and/or bronchoconstriction and/or lung conflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, conflammation, allergies, asthma, impeded respiration, respiratory conflammation, allergies, asthma, impeded respiration, respiratory conflammation, allergies, asthma, impeded respiration, pulmonary transplantation rejection, pulmonary infections, borochitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligosuccent in the target RNA serves to prevent the breakdown of the charget RNA serves to prevent the breakdown of the anti-sense oligos corresponding to the oligosuccent in the target RNA serves to prevent the breakdown of the anti-sense oligos corresponding to the oligosuccent in the target RNA serves to prevent the breakdown of the carget RNA serves to prevent the breakdown of the sense oligos corresponding to the sense oligos corr
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ABD30930
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel composition (a) a first active agent comprising oligonucleotides, effective for alleviating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotides are derived from a gene encoding or regulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bronchoconstriction, respiratory tract inflammation, allers reducing adenosine sensitivity, levels of adenosine (A) or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antialtergic; antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hyportension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human RANTES-derived oligonucleotide SEQ ID 13141.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABD30930 standard;
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                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                  pulmonary transplantation rejection; ss; primer
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24-APR-2001; 2001US-0286036P
                                                                              23-APR-2002; 2002WO-US013143
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             chronic obstructive pulmonary disease; cancer; bronchitis;
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Pred. No. 1.
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ce expression of a target polypeptide associated with lung airway or lung content of a target polypeptide associated with lung airway or lung content on or cancer and can be anti-sense to the corresponding mRNA. Conference invention also describes a kit, that comprises: (a) a delivery content on also describes a kit, that comprises: (b) a delivery content on a santial lergic, anti-inflammatory, antiasthmatic, conference invention has antial lergic, antiinflammatory, antiasthmatic, conference in appointment of the kit. The composition composition is useful for preventing or composition comprises oligo and is administered to reduce the production composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The conduct the amount of target polypeptide present in the lungs. The conduct the admination, allergies and/or bronchoconstriction and/or lung confilammation, allergies, asthma, impeded respiration, respiratory confilammation, allergies, asthma, impeded respiration, respiratory transplantation rejection, pulmonary infections, bronchitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the middless present in the breakdown of the breakdown of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nyce JW, 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies a reducing adenosine sensitivity, levels of adenosine (A) or (A) is surfactant depletion or hyposecretion, when administered to a major oligonucleotides are derived from a gene encoding or regulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted
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                                                                                                   thymidines present in the target RNA serves to prevent the breakdown of
                                                         oligonucleotides into products that free adenosine into , lung, brain, heart, kidney, etc, tissue environment an
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Tang
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L, Shahabuddin
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... or (A) receptors,
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Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension;
beta-adrenergic agonist; respiratory disease; pul-
respiratory distress syndrome; allergic rhinitis;
emphysema; chronic obstructive pulmonary disease;
                                                                                                                                                                                                                                                                                                                                              AI122807-derived
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                                                                                                                                                                                                                                                                                                                                                  oligonucleotide SEQ
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EPIGENESIS

PHARM INC

Homo sapiens.

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating allergies and crediting adenosine sensitivity, levels of adenosine (A) or (A) receptors, comprising adenosine sensitivity, levels of adenosine (A) or (A) receptors, comprises are derived from a gene encoding or regulating composition or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating composition or cancer and can be anti-sense to the corresponding mRNA. Compression of a target polypeptide associated with lung airway or lung composition also describes akit, that comprises: (a) a delivery composition is exparate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, comprises or immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The gulmonary obstruction, and/or bronchoconstriction are associated with a disease or condition such as pulmonary vasoconstriction, compliant and allergies and/or bronchoconstriction are associated with administered pain, cystic fibrosis, allergic rhinitis, pulmonary hypercension, emphysema, chronic obstructive pulmonary disease, pulmonary constructive pulmonary disease, pulmonary to the reduced adenosine content of the arisense oligos corresponding to the oligonucleotides into products that free adenosine into the system e.g., lung, brain, cystic disease, the stream of the content of the content of the service of the content and unwanted effects due to it
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                                                                               Query Match
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Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 15; SEQ ID NO 4421; 763pp; English
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Tang
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L, Shahabuddin
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                                                                                                     1.6e+03;
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ce expression of a target polypeptide associated with lung alivary or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery constituence, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, and peters in the production of the administered composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The composition of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition contains allergies and/or bronchoconstriction and/or lung tinflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as plimonary vascomstriction.

CC inflammation, allergies asthma, impeded respiration, respiratory conditions such as plimonary vascomstriction. The hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system c.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nyce JW,
Miller S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 15; SEQ ID NO 14311; 763pp; English
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                                analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary distriction, respiratory cystic fibrosis, allergic rhinitis, pulmonary
                                                                                                                                                                                                                                                                                                                                                                                            surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, and a carrier and for use of the kit.
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L, Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            on for treating asthma, has antisense ng less percentage of adenosine, targeted to with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         a novel composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            763pp; English.
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emphysema,

chronic obstructive

pulmonary

disease, pulmonary

comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptor surfactant depletion or hyposecretion, when administered to a mammal. I oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The compositions for adding a carrier and for use of the kit.

adding a carrier and has antiallergic, ant

antiinflammatory,

antiasthmatic

The composition

This invention describes a novel composition

(a) a first active agent,

receptors, nammal. The

Claim 15; bronchodilating

SEQ ID

NO 5095;

763pp;

English.

agent.

receptors, nammal. The

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targ nucleic acids associated with lung airway or lung dysfunction

adenosine, targe lung dysfunction,

targeted to

agent,

Miller

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Li Y, Tang

Sandrasagra A, L, Shahabuddin

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Katz

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Pabalan

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Aguilar D;

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24-APR-2001; 2001US-0286036P 23-APR-2002; 2002WO-US013143

(EPIG-) EPIGENESIS PHARM INC

31-OCT-2002 WO200285309-A2 Homo sapiens

WPI; 2003-093058/08

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ABD26083/c
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                                                                                                                                                                                                                                                                                                                                                                                                            respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonuclectides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABD26083 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AA463249-derived oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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RESULT 1412
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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bronchodilating agent.
                        Pharmaceutical composition for treating asthma, oligonucleotide containing less percentage of a nucleic acids associated with lung airway or lu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-OCT-2002
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                                                                                                                                                   WPI; 2003-093058/08
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                                                                                                                                                                                                                                                                                                                                                           24-APR-2001; 2001US-0286036P
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Tang
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L, Shahabuddin
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                           or lung
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                                                          adenosine,
                                                                                      has antisense
                                 dysfunction,
                                                                                                                                                                                                                                   Aguilar D;
                                                             targeted to
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CC surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating CC expression of a target polypeptide associated with lung airway or lung CC dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) constructions for adding a carrier and for use of the kit. The composition CC of the invention has antiallergic, antiinflammatory, antiasthmatic, CC analyssic, hypotensive, immunosuppressive and cytostatic activity, is a CC beta-adrenergic agonist. The composition is useful for preventing or CC treating a respiratory, lung or malignant disease. The administered CC composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to convert the amount of target polypeptide present in the lungs. The composition allocation and/or surfactant humanoraduration and contains allocation and/or surfactant humanoradurations.
                                           inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies reducing adenosine sensitivity, levels of adenosine (A) or (A)
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nt any
            unwanted effects due to it
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á Matches Query Match Best Local 737 GGACTACAGGCGCCCACCAC 756 18; Conser Conservative 1.7%; 0; Score 16.8; Pred. No. 1 Mismatches 6e+03; DB 1; 2 Length 20; Indels 0, Gaps

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2 T; 0 U;

0 Other;

ABD25110/ RESULT 1413 ABD25110; ABD25110 standard; DNA; 20 ВÞ

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GGACTACAGGCGCCCGCTAC 20

AI125228-derived oligonucleotide SEQ ID 4122.

29-JUL-2004

(first entry)

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; 88 primer

Homo sapiens.

WO200285309-A2

2002WO-US013143

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

Claim 15;

SEQ ID

NO 13154; 763pp; English

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RESULT 1414
ABD30932
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Query Match
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Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, all reducing adenosine sensitivity, levels of adenosine (A)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 oligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway of
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                                                analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension;
                                                                                                                                                                                    Human RANTES-derived oligonucleotide SEQ ID 13143.
                                                                                                                                                                                                                       29-JUL-2004
                                                                                                                                                                                                                                                            ABD30932
                                                                                                                                                                                                                                                                                                 ABD30932 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        surfactant depletion or hyposecretion, when administered to a mammal.
                                                                                                          respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic;
                                emphysema; chronic obstructive pulmonary disease; cancer; bronchitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
nes 18; Conserv
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Tang
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                                                                                                                                                 bronchoconstriction; allergy; hyposecretion; pain;
                                                                                                                                                                                                                                                                                                 DNA;
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DT X A X I B B B

29-JUL-2004 ABD32095 ABD32095

(first entry)

밁 S

722 CCTCCTGAGTAGCTGGGACT 741

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CCTCCCGAGTAGCTGGGATT 20

RESULT 1415

standard;

DNA;

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Matches
                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiastimatic, a analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-addrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation allernies and/or eurefactant human-order ton are accordated
                                                                                                                                                                                         inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhintis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) recept surfactant depletion or hyposecretion, when administered to a mammal oligonucleotides are derived from a gene encoding or regulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.
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Tang
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     Gaps
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PDE4C-derived oligonucleotide SEQ ID 14306

bronchoconstriction; allergy;

inflammation;

adenosine

sensitivity; lung; cancer; hyposecretion; pain;

23-APR-2002; 2002WO-US013143 WO200285309-A2 Homo sapiens. emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; surfactant depletion; antiallergic; antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; Miller S, 24-APR-2001; 2001US-0286036P (EPIG-) EPIGENESIS PHARM INC Li Y, Tang ŗ Sandrasagra ndrasagra A, Shahabuddin ŝ Katz E, Pabalan J, Aguilar D;

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

WPI; 2003-093058/08

Claim 15; SEQ ID NO 14306; 763pp; English.

dysfunction or cancer and can be anti-sense to the corresponding mRNN. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasconstriction, and at the attempt increase at the production are associated and the condition and at the sathmatical and activity vasconstriction. inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it surfactant depletion or hyposecretion, when administered to a mammal oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lune comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, aller reducing adenosine sensitivity, levels of adenosine (A) or This invention describes a novel composition (a) a first active agent, allergies (A receptors, or lung

20 BP; 4 Α, 6 C; 7 G; 3 T; 0 U; 0 Other;

Query Match
Best Local Similarity
Matches 18; Conser Conservative 90.0%; 0 Score 16.8; Pred. No. 1 6e+03 DB 1; Length 20; 0; Gaps

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문 S Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; 29-JUL-2004 ABD25776; ABD25776 standard; DNA; pulmonary transplantation rejection; AI085559 DNA fragment 1033 GCTGGGATTACGGGCACCTG GCTGGGATTACAGGCACCCG (first entry) 20 1052 20 ds.

Homo sapiens.

WO200285309-A2

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

Nyce Ä ś Li Y, Sa , Tang L, Sandrasagra A, ,, Shahabuddin Katz S; 'n Pabalan J, Aguilar

WPI; 2003-093058/08

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent. ç

Claim 15; SEQ ID NO 4788; 763pp; English

CC comprising oligonucleotides, effective for alleviating
CC bronchoconstriction, respiratory tract inflammation, alleviating
CC pronchoconstriction, respiratory tract inflammation, alleviating
CC pronchoconstriction, respiratory tract inflammation, alleviating
CC enducting adenosine sensitivity, levels of adenosine (A) or (A) receptors,
CC surfactant depletion or hyposecretion, when administered to a mammal. The
CC oligonucleotides are derived from a gene encoding or regulating
CC expression of a target polypeptide associated with lung airway or lung
CC dysfunction or cancer and can be anti-sense to the corresponding mRNA.
CC The invention also describes a kit, that comprises: (a) a delivery
CC device, in separate containers, (b) the oligonucleotides, (c)
CC instructions for adding a carrier and for use of the kit. The composition
CC of the invention has antiallergic, antiinflammatory, antiasthmatic,
CC analgesic, hypotensive, immunosuppressive and cycostatic activity, is a
Deta-adrenergic agonist. The composition is useful for preventing or
CC treating a respiratory, lung or malignant disease. The administered
CC composition comprises oligo and is administered to reduce the production
CC or availability, or to increase the degradation of the target mRNA or to
CC reduce the amount of target polypeptide present in the lungs. The
CC inflammation allevites and/or bronchoconstriction and/or lung
CC inflammation allevites and/or bronchoconstriction and continue. inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary This invention describes a novel composition (a) a first rejection, active agent,

The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, prevent any unwanted effects due to it

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ABD32092
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        This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung alivay or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20
                                                                                                                                                                                                                                                                     Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200285309-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;
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                                                                                                                                                                                                                                                        bronchodilating agent.
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                                                                                                                                                                                                                        15; SEQ ID NO 14303; 763pp; English.
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, Tang L,
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L, Shahabuddin
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90.0%;
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immunosuppressive and cytostatic
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                                                                                                                                                                                                                                                                                                                                                                                          Pabalan J,
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activity, is a.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC treating a respiratory, lung or malignant disease. The administered condition comprises oligo and is administered to reduce the production cor availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The CC pulmonary obstruction, and/or bronchoconstriction and/or lung cinflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, CC inflammation, allergies, asthma, impeded respiration, respiratory cdistress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary cdistress syndrome, pain, cystic fibrosis, allergic rhinitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system CC eng. lung, brain, heart, kidney, etc, tissue environment and thereby, to crevent any unwanted effects due to it
RESULT 1418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                           Nyce JW,
Miller S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; analgesic; hypotensive;
                                                                                                                                                                                                                                                                                                                                                                                                                                  beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20
                                                                           Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targonucleic acids associated with lung airway or lung dysfunction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human RANTES-derived oligonucleotide SEQ ID 13142.
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                                                                                                                                                                                                                                                         24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                       23-APR-2002; 2002WO-US013143
                                                                                                                                                                                                                                                                                                                       31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                         WO200285309-A2
                                                                                                                                                                                                                                                                                                                                                                                                                       pulmonary transplantation rejection; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABD30931 standard;
                                                              bronchodilating agent.
                                                                                                                                             WPI; 2003-093058/08
                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                           (EPIG-)
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                                                                                                                                                                                                                         EPIGENESIS PHARM INC
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                                                                                                                                                                          Li Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 3 A; 7 C; 4 G; 6 T; 0 U;
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                                                                                                                                                                           Sandrasagra A,
L, Shahabuddin
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                               e of adenosine, targeted or lung dysfunction, and
                                                                                                                                                                                              Pabalan
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This invention describes a novel composition Claim 15; SEQ ID NO 13142; 763pp; English

(a) bi

first active agent,

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RESULT 1419
ABD3103
XX ABD3103
AC ABD3103
XX ADD3103
XX DT 29-JUL-
XX Human;
KW Human;
KW respira
KW surfact
KW surfact
KW surfact
KW emphyse
KW pulmona
XX pulmona
XX Homo sa
XX Homo sa
XX Explora
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CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cyrostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, anlergies and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, condition, such as pulmonary vasoconstriction, respiratory conditions, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human RANTES-derived oligonucleotide SEQ ID 13243.
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                                                                                                                                                                        24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pulmonary transplantation rejection;
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                                                                                          EPIGENESIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ss; primer.
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mammal. The
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Comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating composition, respiratory tract inflammation, allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, considered to a mammal of the coligonucleotides are derived from a gene encoding or regulating considered from a gene encoding or regulating contractant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating considered from a gene encoding or regulating considered from a gene encoding or regulating considered dysfunction of a target polypeptide associated with lung alivary or lung dysfunction of a cancer and can be anti-sense to the corresponding manna. Composition of the invention has antiallergic, antiinflammatory, antiasthmatic, confit the invention has antiallergic, antiinflammatory, antiasthmatic, confit the invention has antiallergic, antiinflammatory, antiasthmatic, composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target manna or to increase the degradation of the target manna or to pulmonary obstruction, and/or bronchoconstriction and/or lung pulmonary obstruction, and/or bronchoconstriction and/or lung pulmonary obstruction, and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, complete manna of the anti-sense oligos corresponding to the middle present in the target manna of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system of the oligonucleotides into products that free adenosine into the system of the oligonucleotides into force the fit of the oligonucleotides into the system of the oligonucleotides of the oligonucleotides of the oligonucleotides of the oligonucleotides of the oligonucleoti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targueleic acids associated with lung airway or lung dysfunction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Miller S,
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                                                       unwanted effects due
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ID NO 13243;
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                                                       to it
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            targeted to
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Matches Query Match Best Local 932 18; Similarity TCACTCTGTTACCCAGGCTG 951 1.7%; 0; Score 16.8; Pred. No. 1. Mismatches 1.6e+03; DB 1; 2 Length 20; Indels 0, Gaps

0

Sequence

20

BP; 3

A; 7 C; 4 G; 6 T; 0 U; 0 Other;

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RESULT 1420
ABD28965
                                       유
ABD28965
                                       ш
standard;
                                      TCACTTTGTCACCCAGGCTG
DNA;
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beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; pulmonary transplantation

N58473-derived oligonucleotide SEQ ID 7977.

(first entry)

Nyce JW,

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Sandrasagra A,

Katz E,

Pabalan

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Aguilar

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RESULT 1421
ABD30414
ID ABD3041
XX
AC ABD3041
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AC ABD3041
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DT 29-JUL-
XX
                                                                                                                                                                                                                                                                                                    CC oligonucleotides are derived from a gene encoding or regulating cc expression of a target polypeptide associated with lung airway or lung conversation or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, considered invention as antiallergic, antiinflammatory, antiasthmatic, considered invention as antiallergic, antiinflammatory, antiasthmatic, considered invention are spiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The composition construction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, confilammation, allergies, asthma, impeded respiration, respiratory construction, pulmonary cyclic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system content and effects due to it
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, aller reducing adenosine sensitivity, levels of adenosine (A) or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleic acids associated bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
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                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            surfactant depletion or hyposecretion, when administered to a mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention describes a novel composition (a) a first active agent,
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                    29-JUL-2004
                                                    ABD30414;
                                                                                   ABD30414 standard;
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                                                                                                                                                                                                                                    l Similarity
18; Conserv
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                                                                                                                                                                                                   TCAAGCAGTCCACCTGCCTC
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Tang
                                                                                                                                                                     TCAAGTAATCCACCTGCCTC 20
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                                                                                                                                                                                                                                                                                                                                       unwanted effects
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                   (first entry)
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L, Shahabuddin S;
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                                                                                     DNA;
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                                                                                                                                                                                                                                                                                                                                     due to it
                                                                                                                                                                                                                                                     Score 16.8;
Pred. No. 1
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                                                                                                                                                                                                                                                                                                      5 T; 0 U;
                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                 Length 20;
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computating origonic respiratory tract inflammation, allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, continued adenosine sensitivity, levels of adenosine (A) or (A) receptors, continued a sensitivity, levels of adenosine (A) or (A) receptors, and contains a gene encoding or regulating continued as are derived from a gene encoding or regulating content on the analyse that comprises: (A) a delivery device, in separate containers, (b) the oligonucleotides, (c) constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antialfammatory, antiasthmatic, canaligesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung pulmonary obstruction, and/or bronchoconstriction and/or lung pulmonary vasoconstriction, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, respiratory can follow the reduced adenosine content of the anti-sense oligos corresponding to the reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system of the oligonucleotides into products that free adenosine into the system of the oligonucleotides of afferts dine to it
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Miller S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; antimonary hypotension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human IL4-R derived oligonucleotide SEQ ID 12625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 15; SEQ ID NO 12625; 763pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-093058/08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 comprising oligonucleotides, effective for alleviating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel composition (a) a first active agent,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          bronchodilating
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, Tang
                                         unwanted effects
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                                           due
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Sequence

20 BP;

5 A;

2 C; 8

9

5 T; 0 U;

0 Other;

Matches Query Match Best Local 18; Similarity Conservative 1.7%; 0; Score 16.8; Pred. No. 1 Mismatches .6e+03; Length Indels 0 Gaps

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29-JUL-2004 ABD26076; ABD26076 standard; (first entry) DNA; 20

₽P

AA463249-derived oligonucleotide SEQ ID 5088

surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human; antisense; respiratory tract bronchoconstriction; allergy; inflammation; adenosine sensitivity; lung; cancer; hyposecretion;

WO200285309-A2

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Li Y, Tang ۲ Sandrasagra A, L, Shahabuddin S Katz E, Pabalan Ģ Aguilar D;

bronchodilating agent. Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and

SEQ ID NO 5088; 763pp; English.

RESULT 1422
ABD26076/c
ID ABD2607
XX
ABD2607 CC comprising oligonucleotides, effective for alleviating
CC treducing adenosine sensitivity, levels of adenosine (A) or (A) receptors,
CC surfactant depletion or hyposecretion, when administered to a mammal. The
CC expression of a target polypeptide associated with lung airway or lung
CC expression of a target polypeptide associated with lung airway or lung
CC dysfunction or cancer and can be anti-sense to the corresponding mRNA.
CC The invention also describes a kit, that comprises: (a) a delivery
CC device, in separate containers, (b) the oligonucleotides, (c)
CC instructions for adding a carrier and for use of the kit. The composition
CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a
CC beta-adrenergic agonist. The composition is useful for preventing or
CC treating a respiratory, lung or malignant disease. The administered
CC composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The
CC uniformation, allergies and/or bronchoconstriction and/or lung
CC inflammation, allergies and/or bronchoconstriction and associated
CC distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary
CC distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary
CC transplantation rejection, pulmonary infections, bronchitis or cancer.
CC The reduced adenosine content of the anti-sense oligon corresponding
CC transplantation content of the anti-sense oligon corresponding
CC transplantation rejection, pulmonary infections, bronchitis or cancer. This invention describes a novel composition (a) a first active agent, adenosine content of the

> ន្តន្តន្តន្ត្ 밁 Ś Query Match Best Local S Matches 18 e.g., lung, prevent any thymidines present in the target RNA serves to prevent the breakdown the oligonuclectides into growcrs that free adenosine into the syste e.g., lung, brain, heart, kidney, etc, tissue environment and thereby prevent any unwanted effects due to it Sequence 20 BP; 8 753 CCACGCCTAGCTAATTITT 772 20 Similarity CCATGCCCAGCTAATTTTTT 1 Conservative A; 2 C; 6 1.7%; <u>.</u> Score 16.8; I Pred. No. 1.6e 0; Mismatches G; 4 T; 0 U; 1.6e+03; 0 Other; DB 1; <u>۷</u> Length 20; Indels 0 Gaps system o f ö 0

ABD26090 standard; DNA; 20 ВÞ

29-JUL-2004 (first entry

AA463249-derived oligonucleotide SEQ ID 5102.

Human; antisense; bronchoconstriction; allergy, hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; pulmonary hyportension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

31-OCT-2002

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

Miller S, Мусе ЈW, Li Y, Sa , Tang L, Sandrasagra A, L, Shahabuddin S Katz 'n Pabalan J, Aguilar

WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, target nucleic acids associated with lung airway or lung dysfunction, bronchodilating agent. targeted ដ

Claim 15; SEQ ID NO 5102; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, suirfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived tride agene encoding or regulating expression of a target polypeptide associated with lung aliway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic. analgesic, hypotensive, beta-adrenergic agonist. immunosuppressive and cytostatic activity, The composition is useful for preventing

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RESULT 1424
ABD26093/c
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antisethmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rininitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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                             Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
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bronchodilating agent
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Tang
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L, Shahabuddin
                                                                                                                                                                                                                                                                                                                PHARM INC
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This invention describes a novel composition (a) a first comprising oligonucleotides, effective for alleviating

active

agent,

Nyce JW, Miller S,

Li Y, Tang

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Sandrasagra A, L, Shahabuddin

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Pabalan

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Aguilar

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SEQ

IJ

NO 5105;

763pp; English

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expression of a target polypeptide associated with lung airway or lung construction or cancer and can be anti-sense to the corresponding mRNA. Conference in the invention also describes a kit, that comprises: (a) a delivery conference, in separate containers, (b) the oligonucleotides, (c) constructions for adding a carrier and for use of the kit. The composition conference in invention has antiallergic, antiinflammatory, antiasthmatic, constantiallergic antiinflammatory, antiasthmatic, constantiallergic antiantimentary, is a beta-adrenargic agonist. The composition is useful for preventing or constaining a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition conference to and/or bornchoconstriction and/or lung confilmmation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasocomstriction.

Construction and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasocomstriction, pain, cystic fibrosis, allergic rhinitis, pulmonary disease, pulmonary administration, respiration, content of the administration content of the administration content of the administration content of the administration cancer.
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RESULT 1425
ABD30936
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                       Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the oligonucleotides into products that free adenosine into e.g., lung, brain, heart, kidney, etc, tissue environment an prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transplantation rejection, pulmonary infections, bronchitis or cancer
The reduced adenosine content of the anti-sense oligos corresponding
thymidines present in the target RNA serves to prevent the breakdown
                                                                                                                                                                                                                                                                                                                                      beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
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                                                                                                                                                                                                                                                                                                                        pulmonary transplantation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human
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                                                                                                      24-APR-2001; 2001US-0286036P
                                                                                                                                                   23-APR-2002; 2002WO-US013143
                                                                                                                                                                                                                                                                              Homo sapiens
                                                              (EPIG-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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to it
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                                                                                                                                                                                                                                                                                                                           primer.
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchits or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic acceptivity, is a management of the containers.
                                                                    Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antisethmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transpiratory.
                                                                                                                                                                                                                                                 Human RANTES-derived oligonucleotide SEQ ID 13246.
                                                                                                                                                                                                                                                                                                                                                                            ABD31035 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim
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ilarity 90.0%;
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sapiens

RESULT 1427 ABD32101

ABD32101 standard; DNA;

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**XXXXXX** 

29-JUL-2004

(first entry)

Human PDE4C-derived oligonucleotide SEQ ID 14312.

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GGCTGGAGTGAAGTGGCACA 20 GGCTGGAGTGCAGTGGCGCA 666 Matches

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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung
                                                                                                                                                                               the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, all reducing adenosine sensitivity, levels of adenosine (A)
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                                                                                                                    Sequence
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                            Score 16.8; DB 1; Pred. No. 1.6e+03;
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CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a CC beta-adrenergic agonist. The composition is useful for preventing or CC treating a respiratory, lung or malignant disease. The administered CC composition comprises oligo and is administered to reduce the production CC or availability, or to increase the degradation of the target mRNA or to CC reduce the amount of target polypeptide present in the lungs. The CC pulmonary obstruction, and/or bronchoconstriction and/or lung CC inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction. CC inflammation, allergies, asthma, impeded respiration, respiratory CC distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation, respection, pulmonary infections, bronchitis or cancer. CC transplantation rejection, pulmonary infections, bronchitis or cancer. CC transplantation rejection, pulmonary infections, bronchitis or cancer. CC thymidines present in the target RNA serves to prevent the breakdown of the oligonuclectides into products that free adenosine into the system.
                                         Query Match
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Miller S,
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                                                                                                                                                                                                                                                 oligonucleotides into products that ., lung, brain, heart, kidney, etc, t vent any unwanted effects due to it
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                                                                                                                                                                        BP; 3 A; 5 C; 6 G; 6 T; 0 U; 0 Other;
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L, Shahabuddin S;
                                         1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      763pp; English
Score 16.8; D
Pred. No. 1.6e
O; Mismatches
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                                                                                                                                                                                                                                                                                                   tissue environment and thereby,
                                                                                      DB 1; Length 20;
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791

GGGGTTCACCATGTTCGCCA 810

Matches

Similarity

0,

1.6e+03;

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Gaps

0

RESULT 1428
ABD30945
ID ABD30945
XX ABD30945
AC ABD30945
AC ABD30944
XX ABD30945
AC ABD3094
AC ABPRAC ABBANA
AC ABBANA comprising oligonucleotides, effective for alleviating
CC comprising oligonucleotides, effective for alleviating
CC control of the control of **A** Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. ABD30945 standard; Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and Human RANTES-derived oligonucleotide SEQ ABD30945; Miller 23-APR-2002; 2002WO-US013143 29-JUL-2004 Claim 15; SEQ ID NO 13156; 763pp; English WPI; 2003-093058/08 24-APR-2001; 2001US-0286036P WO200285309-A2 Homo sapiens. This invention describes a novel composition (a) (EPIG-) EPIGENESIS PHARM INC ß GGGTTTCACCATGTTGGCCA 20 Li Y, Tang (first entry) Sandrasagra A, DNA; Shahabuddin 20 ŝ Katz E, ID 13156 Pabalan a first active agent, 4

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e.g.

oligonucleotides into products that free adenosine into the system ., lung, brain, heart, kidney, etc, tissue environment and thereby, vent any unwanted effects due to it

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Matches
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surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung alrway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic aggnist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                           This invention describes a novel composition (a) a first active age comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) recompositions of the control of the c
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PDE4C-derived
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Tang
                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 14341;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sandrasagra A,
., Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 763pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
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                                                                                                                                                                                                                                                                                    (A) receptors,
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inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                   composition comprises oligo and is administered to reduce the or availability, or to increase the degradation of the target reduce the amount of target polypeptide present in the lungs pulmonary obstruction, and/or bronchoconstriction and/or lung
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BP; 4 A;
  (C)
  ω
  ຸດ
  7 T;
0 U;
  0 Other;
                                                                                                                                                                                                                                                                                                                                                                          production mRNA or to
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Query Match Best Local S Matches 18 18; Similarity Conservative 1.7%; 90.0%; 0 Score 16.8; Pred. No. 1 Mismatches 1.6e+03 DB 1; Length Indels 0 Gaps

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밁 δ 751 CACCACGCCTAGCTAATTTT 770 1 CACCATGCCTGGCTAATTTT 20

RESULT 1430 ABD28960

ABD28960 standard; DNA;

20

N58473-derived oligonucleotide SEQ ID 7972.

(first entry)

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hyportension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

Nyce JW, Miller S, Li . Tang ) ŗ Sandrasagra A, , Shahabuddin s Katz Ή Pabalan 'n Aguilar

WPI; 2003-093058/08

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targenucleic acids associated with lung airway or lung dysfunction bronchodilating agent. dysfunction, ç

Claim 15; SEQ ID NO 7972; 763pp; English

This invention describes a novel composition (a) a first active comprising oligonucleotides, effective respiratory tract for alleviating inflammation, a

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RESULT 1431
ABD26095/c
CC analgesic, hypotensive, immonsuppressive and cytostatic activity, is a cc beta-adrenergic agonist. The composition is useful for preventing or cc treating a respiratory, lung or malignant disease. The administered cc composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to creduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung cc inflammation, allergies and/or surfactant hypoproduction are associated cwith a disease or condition such as pulmonary vasoconstriction, cc inflammation, allergies, asthma, impeded respiration, respiratory cc distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system ce.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antisethmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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                                 Nyce JW,
Miller S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AA463249-derived oligonucleotide SEQ ID 5107.
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                                                                                                                                                                                                                                       24-APR-2001; 2001US-0286036P
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                                                                                                                                                           (EPIG-) EPIGENESIS PHARM INC
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    Li ..
Tang /
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                                     Sandrasagra A,
,, Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ..
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Pred. No. 1
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                                                                           Katz
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                                                                           Pabalan
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                                                                               Aguilar
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ce expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. Comprises: (a) a delivery constitution or cancer and can be anti-sense to the corresponding mRNA. Comprises: (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition confirments of the invention has antiallergic, antiinflammatory, antiasthmatic, composition has antiallergic, antiinflammatory, antiasthmatic, composition are associated composition comprises oligo and is administered to reduce the production corresponding or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, allergies and/or bronchoconstriction and/or lung confilammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vascoconstriction, distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the charget and or that free adenosine into the arget RNA serves to prevent the breakdown of the pulmonary that are the production of the arget and not the production of the arget that free adenosine into the arget RNA serves to prevent the breakdown of the production in the target RNA serves to prevent the breakdown of the production in the production of the arget RNA serves to prevent the breakdown of the production in the production i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             surfactant depletion or hyposecretion, when administered to a mooligonucleotides are derived from a gene encoding or regulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allers reducing adenosine sensitivity, levels of adenosine (A) or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bronchodilating agent.
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                                                                                                                   the oligonuc.
                                                                  brain, heart,
      unwanted
leotides into products that free adenosine into
brain, heart, kidney, etc, tissue environment ar
unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               БЛ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        receptors,
mammal. The
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밁 S Matches Query Match Local 472 AGGATGAAGTGCAGTGGTGT 491 18; Similarity Conservative 1.7%; .. Pred. Score 16.8; Pred. No. 1 Mismatches 1.6e+03; 2; DB 1; Length 20 Indels 0; Gaps

0

Sequence

20

BP;

ហ A; 8

C; 3

G; 4 T; 0 U;

0 Other;

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ABD32089 standard; DNA; 20 20 AGCCTGAAGTGCAGTGGTGT 1

Human PDB4C-derived oligonucleotide 29-JUL-2004 (first entry) SEQ ID 14300.

ABD32089;

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antialflammatory; artisthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction pulmonary transplantation rejection; ss; respiratory distress syndrome; allergic rhinitis; emphysema; chronic obstructive pulmonary disease; ratory disease; pulmonary vasoconstriction; allergic rhinitis; pulmonary hypertension; primer. cancer; bronchitis;

WO200285309-A2

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                                                                                                                                                                                                                                                                                                                                                                                                              CC comprising oligonuclectides, effective for alleviating composition of ligonuclectides, effective for alleviating compositions sensitivity, levels of adenosine (A) or (A) receptors, composition or hyposecretion, when administered to a mammal. The coligonuclectides are derived from a gene encoding or regulating contribution of a target polypeptide associated with lung airway or lung composition of a target polypeptide associated with lung airway or lung consistent in the invention also describes a kit, that comprises: (a) a delivery consistent in the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cyrostatic activity, is a composition comprises oligo and is administered to reduce the production composition, allergies and/or badinistered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, allergies, asthma, impeded respiration, respiratory conditions, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation, allergies, asthma, impeded respiration, respiratory conditions polyperide present in the lungs. The conformation, allergies and/or surfactant hypoproduction are associated conformation, allergies, asthma, impeded respiration, respiratory conditions pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonary brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                 S
                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 18
                     Human PDE4C-derived oligonucleotide SEQ ID 14330.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
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                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 15; SEQ ID NO 14300; 763pp; English
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                                                           29-JUL-2004
                                                                                                                                    ABD32119 standard; DNA; 20
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Tang
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                                                                                                                                    ВP
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                                                                                                                                                                                                                                                                                                                           Score 16.8; DB 1;
Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                            DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                  0 Other;
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646 AGGCTGGAGTGCAGTGGCGC 665

AGGCTGGAGTGCAGTGATGC

Matches Query Match Best Local &

18;

Conservative

<u>.</u>

Mismatches

Indels

<u>,</u>

Gaps

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20

Local Similarity

1.7%;

Score 16.8; Pred. No. 1.

1.6e+03

Sequence

20

BP; 4

A; 3 C; 9 G; 4 T; 0 U; 0 Other;

prevent any unwanted

effects due

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Consider the content of the composition of the target polypeptide associated with lung alivary or lung content of a target polypeptide associated with lung alivary or lung content of a target polypeptide associated with lung alivary or lung content of the invention or cancer and can be anti-sense to the corresponding mRNA.

Constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analysis, antiasthmatic, and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, allergies and/or bronchconstriction and/or lung lunguary obstruction, allergies and/or bronchconstriction and/or lung conflammation, allergies and/or bronchconstriction, and sease or condition such as pulmonary vasoconstriction, conflammation, allergies, asthma, impeded respirator, respiratory hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary conflammation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system content of the areaset NAA serves to prevent the breakdown of the oligonucleotides into afforts due to the products and thereby, to respect the content of afforts due to the system content and unserved defense and the content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system content of a feforts due to the system content of the anti-sense environment and thereby, to
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Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   respiratory tract inflammation; adenosine sensitivity; lung; carcer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) recept surfactant depletion or hyposecretion, when administered to a mammal. oligonucleotides are derived from a gene encoding or regulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targueleic acids associated with lung airway or lung dysfunction
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes a novel composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bronchodilating agent.
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, Tang L,
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L, Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   dysfunction,
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RESULT 1434
ADF86417
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ADG86786
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DXXXXXXXXXXXXXXXX
                                                                                                                                                                                                                                                                     Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                       candidates. The present DN used in an example of the
                                                                                                                                                                                                                                                                                                                                                  antagonists of the invention may be used to treat, prevent and diagnose acute leukaemia, the VLA4 antagonists may also be used to screen drug candidates. The present DNA sequence represents a PCR primer that was
                                                                                                                                                                                                                                                                                                                                                                                                                                          Treatment and/or prevention of acute leukemia with medicinal compositions containing VLA4 antagonist, also applicable in diagnosing its prognosis and screening drug candidates.
                                                                                                                                                                                                                                                                                                                                                                                  The invention comprises VLA4 antagonists that may optionally be used with other anticancer agents for the treatment of acute leukaemia. The VLA4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-FEB-2004
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                                          Human; ss; PPAR delta; peroxisome proliferative activated receptor delta; antisense gene therapy; cytostatic; osteopathic; antidiabetic; cancer; osteoporosis; diabetes; endocrine disorder.
                                                                                                                 11-MAR-2004
                                                                                                                                                              ADG86786 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                  Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                        Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-012487/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Niitsu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (NIIT/) NIITSU Y.
(MATS/) MATSUNAGA T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-MAY-2002; 2002WO-JP004704
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-MAY-2002; 2002WO-JP004704
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-NOV-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VLA4 antagonist; acute leukaemia; screening; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VLA4 antagonist-related PCR primer #2
                                                                                         Human
                       Homo sapiens.
                                                                                                                                                                                                                                             663
                                                                                                                                                                                                                                                                    18;
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                                                                                                                                                                                                                                     CGCAATCTTGGCTCACTGCA 682
                                                                                                                                                                                                                      CGCGATCTCGGCTCACTGCA 20
                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Takemoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matsunaga T,
                                                                                         antisense
                                                                                                                                                                                                                                                                                                                  BP; 3 A;
                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                               1.7%;
                                                                                                                                                                                                                                                                                                                  8 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                        oligonuclotide
                                                                                                                                                                                                                                                                                                                                                                                                                      72pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Miyake K,
                                                                                                                                                                                                                                                                                                                                         invention.
                                                                                                                                                               ВP
                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                Score 16.8;
Pred. No. 1
                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sakamaki S,
                                                                                           ISIS 136865
                                                                                                                                                                                                                                                                                 1.6e+03
                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                     <u>ب</u>
                                                                                                                                                                                                                                                                                           Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Akiyama
                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ħ,
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                                                                                                                                                                                                                                                                     Gaps
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Key

Location/Qualifiers

Human; ds; PPAR delta; peroxisome proliferative activated receptor delta;

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SXEXEXEX B
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                                                                                                                          RESULT 1436
ADG86939/c
                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                      of a preferred target region on a nucleic acid molecule encoding PPAR-delta and a composition comprising the antisense oligonucleotide and a carrier. The antisense oligonucleotide comprises at least one modified internucleoside linkage (preferably a phosphorothioate linkage), at least one sugar moiety (preferably 2'-O-methoxyethyl moiety) and at least one modified nucleobase (which is a 5-methyl cytosine). The antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an antisense oligonucleotide comprising 8-80 nucleobases in length targeted to the coding region of a nucleic acid molecule encoding PRAR-delta (peroxisome proliferative activated receptor delta), where the antisense compound inhibits the expression of the PPAR-delta and has any of the 66 sequences of 20 amino acids fully defined in the specification. Also included are a compound of 8-80 nucleobases in length that specifically hybridises with at least an 8-nucleobase portion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>.</u>
                                                                                                                                                                                                                                                                                                                             compounds are useful for treating cancer, osteoporosis, diabetes or various endocrine disorders. The Human PPAR delta gene is located on chromosome 6521. The present sequence is an antisense oligonucleotide the invention targeting human PPAR delta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides of 8-80 nucleobases, useful for treating cancer, diabetes, osteoporosis or various endocrine disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-022078/02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaarde W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-MAY-2002; 2002US-00160807
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                                                                  Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2003224514-A1
                          Human PPAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC
                                                       11-MAR-2004
                                                                                  ADG86939;
                                                                                                             ADG86939 standard; DNA; 20
                                                                                                                                                                                                                1027 CAAGCAGCTGGGATTACGGG 1046
                                                                                                                                                                                                                                            18;
                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Freier SM,
                          antisense oligonucleotoide target sequence #1.
                                                                                                                                                                                                                                                                                                  BP; 6 A; 3 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2002US-00160807
                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag= C
/mod_base= OTHER
/note= "2'-methoxyethyl residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag=
/mod_ba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "Phosphorothioate linkages and all cytidines are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= b
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              base=
                                                                                                                                                                                                                                                          90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         155pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AT;
                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                       Score 16.8;
                                                                                                                                                                                                                                                           Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            English.
                                                                                                                                                                                     20
                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                          No.
                                                                                                                                                                                                                                                          .6e+03
                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                      Length 20;
                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                         The antisense sis, diabetes or
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                                                                                                                                                                                                                                              Gaps
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ADH56917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                delta), where the antisense compound inhibits the expression of the PPAR-
delta and has any of the 66 sequences of 20 amino acids fully defined in
the specification. Also included are a compound of 8-80 nucleobases in
length that specifically hybridises with at least an 8-nucleobase portion
of a preferred target region on a nucleic acid molecule encoding PPAR-
delta and a composition comprising the antisense oligonucleotide and a
carrier. The antisense oligonucleotide comprises at least one modified
internucleoside linkage (preferably a phosphorothioate linkage), at least
one sugar mojety (preferably 2'-0-methoxyethyl mojety) and at least one
modified nucleobase (which is a 5-methyl cycosine). The antisense
compounds are useful for treating cancer, osteoporosis, diabetes or
various endocrine disorders. The Human PPAR delta gene is located on
chromosome 6p21. The present sequence is a human pPAR delta genomic
target sequence for the antisense oligonucleotides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to an antisense oligonucleotide comprising 8-80 nucleobases in length targeted to the coding region of a nucleic acid molecule encoding PPAR-delta (peroxisome proliferative activated receptor molecule encoding proliferative activated receptor molecule encoding proliferative activated receptor molecule encoding proliferative encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense oligonucleotides of 8-80 nucleobases, useful for treating cancer, diabetes, osteoporosis or various endocrine disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-022078/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaarde W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAY-2002; 2002US-00160807.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-MAY-2002; 2002US-00160807.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US2003224514-A1
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                                                                                            Homo sapiens.
                             US2003219810-A1
                                                                                                                                                                                  inflammation; chronic obstructive pulmonary disease;
rheumatoid arthritis; inflammatory bowel; psoriasis; asthma;
antiasthmatic; antiinflammatory; antiallergic; pharmacogenomic; forensic;
                                                                                                                                                                                                                                                                                                                                                   Human CARD4 DNA oligo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                       paternity testing;
                                                                                                                                                                                                                                                                                     ss; human; CARD4; NOD1; CED4/Apaf-1; caspase-9 induced apoptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADH56917 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1027 CAAGCAGCTGGGATTACGGG 1046
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 4 A; 7 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 175; 155pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                       single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.7%;
                                                                                                                                                                                                                                                                                                                                                comprising an allelic variant SeqID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Watt
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 16.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                                                                                                                                                       polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 20;
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RESULT 1438
ADH73294/c
ID ADH7329
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    arthritis, inflammatory bowel disease, psoriasis or asthma. According the compositions of this invention exhibit antiasthmatic, antiinflammatory and antiallergic activities. Furthermore, they may be used to identify patients that would be strong candidates for effective treatment with a CARD4 modulator, in pharmacogenomics, or in monitorist the effects of CARD4 therapeutics during clinical trials. The nucleic acid molecule may also be used in forensics or paternity testing. This oligonucleotide sequence is a human CARD4 DNA oligo comprising an allegature.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention relates to novel single nucleotide polymorphisms within the human CARD4 gene. Specifically, it refers to allelic variants of CARD4 (NODI), a member of the CED4/Apaf-1 family that is involved in caspase-9 induced apoptosis and inflammation. The present invention describes a kit for determining the allelic variants of CARD4 polymorphic regions of an individual, which can be useful for predicting susceptibility, as well as diagnosis, prevention and treatment of various displaying including chronic obstructive pulmonary disease, rheumatoid districtive including chronic obstructive pulmonary disease, rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BARN/)
(BERT/)
                                                                                                                                                                                          epigenetic abnormality detection; hypomethylated sequence; multi-copy DNA element; retroelement; Alu sequence; Huntin schizophrenia; bipolar disorder; human; PCR; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated nucleic acid molecule comprising an allelic variant of a CARD4 gene, useful for diagnosing, preventing or treating asthma or a apoptotic, inflammatory or allergic disorder, or in pharmacogenomics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Barnes G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-MAR-2002; 2002US-0368184P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-MAR-2003; 2003US-00401194
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                              06-JUN-2002; 2002US-0386818P
                                                              06-JUN-2003; 2003WO-CA000820
                                                                                                                                                                                                                                                          Human Alu sequence
                                                                                                                                                                                                                                                                                            25-MAR-2004 (first entry)
                                                                                                                                                                                                                                                                                                                            ADH73294;
                                                                                                                                                                                                                                                                                                                                                             ADH73294 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-010870/01.
                                                                                              18-DEC-2003
                                                                                                                               WO2003104487-A2
                                                                                                                                                             Homo sapiens.
(ADDI-) CENT ADDICTION & MENTAL HEALTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
mes 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         384 CTCCCAAAGTGCTGGGATTA 403
                                                                                                                                                                                                                                                                                                                                                                                                                                            ۲
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                            CTCCCAAAGCACTGGGATTA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bertin J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                          PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.7%;
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                                                                                                                                                                                                                                                                                                                                                             BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.8; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                 ; Alu .
PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.6e+03
                                                                                                                                                                                             sequence; Huntington's
R; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        comprising an allelic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for effective in monitoring
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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RESULT 1439
ADI30044
ID ADI3004
XX ADI3004
XX ADI3004
XX Human d
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             retroelement - endogenous retroviral sequences (ERV), SINE sequences, Alu sequences, LINE sequences and LI sequences. The method of the invention is useful for detecting a genetic abnormality associated with a disease, e.g. Huntington's disease, schizophrenia or bipolar disorder. The present DNA sequence represents a human Alu sequence PCR primer that was used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention comprises a method for detecting an epigenetic abnormality associated with a disease. The method involves identifying, within a eukaryotic genome, a locus having a hypomethylated sequence specific for the disease and an endogenous multi-copy DNA element, such as a retroalement - endogenous retroviral sequences (ERV), SINE sequences, Alu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-062375/06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADI30044;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                            18-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Antisense therapy; human; dual specific phosphatase 4; hyperproliferative disorder; developmental disorder; apoptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADI30044
                                                                                                                                                                                                                                                                             17-JUN-2002; 2002US-00174460
                                                                                                                                                                                                                                                                                                                                                17-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            389
                                               2004-061286/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     dual specific phosphatase 4 DNA, antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20
                                                                                                                          ₿₽,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                  SISI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAACTGCTGGGAGTACAGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAAGTGCTGGGATTACAGGC 408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of the invention.
                                                                                                                          Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 3 A; 7 C; 4 G; 6 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                    2002US-00174460
                                                                                                                                                                                                      PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note = "This oligonucleotide has a phosphorothioate backbone and 2'.methyoxyethyl (2'.MOE) wings at the backbone should be not be not benefit at early a nucleotides in length at early and 3' ends, which are 5 nucleotides in length at early and 1'. The nucleotides in length at length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2; 257pp; English
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                                                                                                                          Dobie
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Pred. No. 1.6e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 #64
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FEFFXXXSSSSSSSSSSSSSSSSS
                                                                                                                                                                            New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding dual specific phosphatase 4, useful for treating cancer, developmental disorder or a condition arising from aberrant
                                                                                                                                                                       apoptosis.
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SEQ Ü NO 77; 61pp; English

nucleic acid encoding dual specific phosphatase 4. The antisense compound comprises an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibits the expression of dual specific phosphatase 4. The antisense oligonucleotide is a chimeric oligonucleotide. The antisense oligonucleotide comprises at least one modified internucleoside linkage, preferably a phosphorothicate linkage. It also comprises at least one modified sugar moiety, preferably a 2'-O-methoxyethyl (2'-MOE) sugar moiety. The antisense oligonucleotide further comprises at least one modified nucleobase, preferably a 5-methyloytosine. The antisense oligonucleotides are useful for the treatment of diseases such as hopermynliferative discontained. hyperproliferative disorders, developmental disorders, and diseases associated with aberrant apoptosis. The present sequence represents an antisense oligonucleotide used in the examples of the present invention The present invention relates to antisense compounds targeted nucleic acid encoding dual specific phosphatase 4. The antiser ö invention

Sequence 20 BP; 2 A; 7 C; 4 G; 7 T; 0 U; 0 Other;

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           á
                          Matches
                                Query Match
Best Local S
              836 TGATCTGCCTGCCTCGGCCT 855
                           18;
\vdash
                                  Similarity
TGATCTGCCTGCCTCAGTCT
                          Conservative
                                  1.7%;
                           0
                                       Score
                                  Pred.
  20
                            Mismatches
                                   16.8;
No. 1
                                  No.
                                  .6e+03
                                         DB
                                         -
                            2;
                                         Length
                             Indels
                                         20
                            0
                             Gaps
                             0
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RESULT 1440
ADH76711
ID ADH7671
XX ADH7671
XX ADH7671
XX PAPR-
XX MCHR1 9
XX MCHR1 9
XX Unident
XX Unident
XX WO20031
XX Unident
XX Unident
XX Unident
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XX Unident
XX WO20031
XX Unident
XX WO20031
XX Unident
XX WO18-DEC-
XX WO20031
XX Unident
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XX Unident
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XX Copplyee
CC Coppabl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADH76711;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADH76711 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       analysis primer #20
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obesity; primer; melanin-concentrating hormone receptor 1; MCHR1; anorectic; 88 gene therapy

Unidentified

WO2003104489-A2

18-DEC-2003

05-JUN-2003; 2003WO-EP005917

05-JUN-2002; 2002EP-00012569

(UYPH-) UNIV PHILIPPS MARBURG

Platzer M, Reichwald Platzer C, Gudermann 'n Hebebrand ç Hinney

WPI; 2004-062377/06.

New diagnostic composition, useful presence of a molecular variant of the disorder. for the diagnosing obesity related to the MCHR1 gene or a susceptibility to

Example 2; Page 42; 76pp; English.

The polynucleotide composition comprises: a sequence encoding polypeptide with defined sequences given in the specification; capable of hybridizing to a melanin-concentrating hormone rece invention relates to a novel diagnostic polynucleotide composition. a sequence

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RESULT 1441
ADH76713/c
ID ADH7671
XX MCHR1 9
XX MCHR1 9
XX Mobesity
XX M
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Reichwald I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MCHR1) gene; a polynucleotide encoding an MCHR1 polypeptide; or a sequence comprising one or more of the nucleotide exchanges (SNP's) given in the specification and at least 8 bases of surrounding sequence of the MCHR1 gene. The composition has anorectic activity. The polynucleotide composition may be used in gene therapy to treat the disorders of the invention. The composition is useful for diagnosing obesity related to the presence of a molecular variant of MCHR1 gene or a susceptibility to the disorder. The MCHR1 protein or polynucleotide is useful for preparing a medicament for treating or preventing obesity related to the presence of a molecular variant of the MCHR1 gene. This polynucleotide
The polynucleotide composition comprises: a sequence encoding a polypeptide with defined sequences given in the specification; a sequence capable of hybridizing to a melanin-concentrating hormone receptor 1 (MCHRI) gene; a polynucleotide encoding an MCHRI polypeptide; or a sequence comprising one or more of the nucleotide exchanges (SNP's) given in the specification and at least 8 bases of surrounding sequence of the MCHRI gene. The composition has anorectic activity. The polynucleotide composition may be used in gene therapy to treat the disorders of the invention. The composition is useful for diagnosing obesity related to the disorder. The MCHRI protein or polynucleotide is useful for presence of a molecular variant of the MCHRI gene or a susceptibility to the disorder. The MCHRI protein or polynucleotide is useful for treating or preventing obesity related to the presence of a molecular variant of the MCHRI gene. This polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-062377/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-JUN-2002; 2002EP-00012569
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JUN-2003; 2003WO-EP005917
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     obesity; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                melanin-concentrating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MCHR1 genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 42; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a novel diagnostic polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYPH-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diagnostic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    UNIV PHILIPPS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of a molecular variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GTGCAGTGGTGATCTCGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence analysis primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         composition,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
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riant of
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20
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MCHR1 gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2
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                                                                                                                                                                                                                                                                                                                                                                                                                            composition.
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RESULT 1442
ADH77272/c
ID ADH77272 standard; DNA;
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Best Local (
      The invention describes a compound 8-80 nucleobases in length targeted to, and which specifically hybridises with a nucleic acid molecule encoding a PAZ/PIWI domain-containing protein, and inhibits the expression of a PAZ/PIWI domain-containing protein. The compound, composition and methods are useful for treating a disease or condition associated with PAZ/PIWI domain-containing protein, such as a associated with PAZ/PIWI domain-containing protein, such as a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cytostatic; PAZ/PIWI domain-containing protein inhibitor; PAZ/PIWI domain-containing protein; hyperproliferative disorder; cancer; aberrant cellular differentiation; human; PAZ/PIWI domain-containing protein; antisense technology; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   represents an MCHR1 primer of the invention.
                                                                                                                          New antisense oligonucleotide targeted to a nucleic acid encoding PAZ/PIWI domain-containing protein, useful for treating cancer or disease arising from aberrant cellular differentiation.
                                                                                                                                                                               WPI; 2004-052174/05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PAZ/PIWI domain-containing protein oligo
                                                                                                  Example 15;
                                                                                                                                                                                                                                (ISIS-) ISIS
                                                                                                                                                                                                                                                        17-JUN-2002;
                                                                                                                                                                                                                                                                                  17-JUN-2002;
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                                                                                                 SEQ ID NO 162; 119pp; English
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                                                                                                                                                                                                                                 PHARM INC
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/note= "OTHER= Phosphorothioate backbone. All cytidine
residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                              note=
                                                                                                                                                                                                                                                                                                                                                                           mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                           note= "OTHER= 2'-O-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                       *tag= c
                                                                                                                                                                                                                                                                                                                                                                                                                              mod base=
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90.0%;
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                               (2'-MOE)
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Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    arising from aberrant cellular differentiation. They are also useful in research and diagnostics for modulating the expression of PAZ/PIWI domacontaining protein. This sequence represents a human PAZ/PIWI domain-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cytostatic; PAZ/PIWI domain-containing protein inhibitor; PAZ/PIWI domain-containing protein; hyperproliferative diaberrant cellular differentiation; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human PAZ/PIWI domain-containing protein oligo segid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADH77198;
                                                                              New antisense oligonucleotide targeted to a nucleic aciopAZ/PIWI domain-containing protein, useful for treating disease arising from aberrant cellular differentiation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PAZ/PIWI domain-containing protein; antisense antisense oligonucleotide; ss.
The invention describes a compound 8-80 nucleobases in length targeted to, and which specifically hybridises with a nucleic acid molecule encoding a PAZ/PIWI domain-containing protein, and inhibits the expression of a PAZ/FIWI domain-containing protein. The compound,
                                                                                                                           WPI; 2004-052174/05
                                                                                                                                                                                                17-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                          SEQ ID NO 88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
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                                                                                                                                                                          PHARM
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                                                                                                                                                                                                                                                                                                                                                                                     residues are
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/note= "OTHER= 2'
                                                                                                                                                                                                                                                                                                                                                                                    note= "OTHER= Phosphorothioate residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
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                                                          119pp; English.
                                                                                                                                                                                                                                                                                           OTHER
HER= 2'
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                                                                                                                                                                                                                                                                                                                                         -O-methoxyethyl
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No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                backbone.
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                                                                                                                                                                                                                                                                                            (2'-MOE)
                                                                                                                                                                                                                                                                                                                                         (2'-MOE)
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                                                                                                       acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disorder;
                                                                                            d encoding cancer or
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CGCAAAGTGCTGGGATGACA 1 CCCAAAGTGCTGGGATTACA 405 Query Match Best Local S Matches 18

Similarity

90.0%;

Score 16.8; Pred.

Conservative

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Mismatches

Indels

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RESULT 1444
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Matches 18
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                                                                                 The invention comprises antisense oligonucleotides that are targeted to nucleic acid encoding P2X4. The antisense oligonucleotides are useful for inhibiting the expression of P2X4 in cells or tissues to treat diseases associated with P2X4 expression, such as: neurological disorders, bone disorders (e.g. osteoporosis), or rheumatoid arthritis. The present DNA sequence represents an antisense oligonucleotide for the human P2X4 gene The present DNA sequence is a 2'-O-methoxyethyl gapmer with a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense oligonucleotide; P2X4; P2X4-associated diseases; neurological disorder; bone disorder; osteoporosis; rheumatoid arthritis; 2'-O-methoxyethyl gapmer; phosphorothioate backbone; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotides for modulating P2X4 expression, diagnosing, preventing or treating conditions associated with ineurological disorders, osteoporosis or rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JUL-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUL-2002; 2002US-00187659.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADI81381 standard; DNA;
      Sequence
                                                                                                                                                                                                                                                                                                                         Example 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-081656/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2004002152-A1
                                                         The present DNA sequence i
phosphorothioate backbone.
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      BP;
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                                                                                                                                                                                                                                                                                                                         SEQ ID NO 13; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PHARM INC
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      3 A;
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      7 C; 4
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         G; 6 T; 0 U; 0 Other;
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Pred. No.
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No. 1
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RESULT 1445
ADJ36817
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Best Local (
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                                                                                          The invention describes a new isolated nucleic acid comprising a fully defined sequence having 23574 bp or at least its 50 or 15 contiguous nucleotides and includes: allele G of single nucleotide polymorphism (SNP) AB+2; allele G of SNP BC+1; and allele C of SNP BC+2. The invention describes identifying increased susceptibility to a disorder comprising asthma, bronchial hyperresponsiveness, atopy, chronic obstructive lung disease and adult respiratory distress syndrome in a subject comprising testing a biological sample obtained from a subject for the presence of at least one allele or haplotype given in the specification, where the presence identifies an increased susceptibility to the disorder. The nucleic acid is useful for preparing a composition for treating disorderse comprising asthma, bronchial hyperresponsiveness, atopy, chronic obstructive lung disease and adult respiratory distress syndrome. This sequence represents a primer used to detect single nucleotide
                                                       Seguence
                                                                                 sequence represents a primer used to polymorphisms in the human gene 216.
                                                                                                                                                                                                                                                                                                         Example 10; SEQ ID NO 208; 441pp; English.
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13-APR-2001; 2001US-00834597.
19-APR-2002; 2002US-00126022.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human gene 216 SNP
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DEL MASTRO R
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                                                                                                                                                                                                                                                                                                                                    nucleic acid, useful for preparing a composition for tre
s e.g., asthma, bronchial hyperresponsiveness, atopy, chr
ive lung disease and adult respiratory distress syndrome.
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Allen K, P
 Conservative
                                                      BP; 4
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             Score 16.8; DB 1
Pred. No. 1.6e+03
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RESULT 1447 ADJ59777

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RESULT 1446
ADJ60954
ID ADJ6095
XX ADJ6095
XX Oligonu
XX Oligonu
XX interle
KW airway
KW cystic
KW pulmona
XX Pom volumona
XX Novel (EPIG-)
XX Pom volumona
XX Novel (EPIG-)
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant generical, RANTES, MCP4, useful for prophylaxis or treating respiratory
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                                                                                                                                                                                                                                                                                                           Sequence 20
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                                                                            179
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AGTAGAGATGGGGTTTCACC
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Pred. No. 1.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nyce JW, Tang
Shahabuddin S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo
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                                                                                                                                                                                                                                                                                                Sequence
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       06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                   obstruction.
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                                                                        ADJ59866 standard;
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                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                              GGACTACAGGCGCCCACCAC 756
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                                                                                                                                                            GGACTACAGGCGCCCGCTAC
                                                                                                                                                                                                                                                                                                BP; 4 A;
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Lu
     (first entry)
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                                                                                                                                                                                                                                                                                                                                                   present sequence
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H, Cong H;
                                                                          DNA;
                                                                                                                                                                                                                                           1.7%;
                                                                                                                                                                                                                                                                                                8 C; 6 G; 2 T; 0 U; 0 Other;
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Pred. No. 1
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                                                                                                                                                                                                                                               1.6e+03;
                                                                                                                                                                                                                                                               DB 1;
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                                                                                                                                                                                                                                                               Length 20;
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RESULT 1449
ADJ60948
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Cc interleukin (IL)-4 receptor, IL-5 receptor or salts of the coligonucleotide and optionally surfactant operatively linked to the coligonucleotide. The method is useful for preventing or treating a cc respiratory or lung disease, which involves administering to the airways cf a subject an effective amount of an inhibitor. The oligonucleotide is cc useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded cc respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (CRDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide associated to RANTES #115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nyce JW, Tang L, San Shahabuddin S, Lu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-FEB-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-203534/19.
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interleukin; IL-4 receptor; IL-5 receptor; lur airway inflammation; allergy; asthma; impeded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20
                                                                                    Oligonucleotide associated
                                                                                                                                              06-MAY-2004
                                                                                                                                                                                                                                                    ADJ60948 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                  932 TCACTCTGTTACCCAGGCTG 951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2; SEQ ID NO 722;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; 7 C; 4 G; 6 T; 0 U; 0 Other;
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                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          85pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 16.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. 1.6e+03
                                                                                          PDE4C #14.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
           lung disease;
ded respiration;
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                                        disease;
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RESULT 1450
ADJ59764
ID ADJ5976
XX ADJ5976
XC ADJ5977
XC ADJ597
XC 
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases
                                                                                                              interleukin; IL-4 receptor; IL-5 receptor; lung disease;
airway inflammation; allergy; asthma; impeded respiration;
cystic fibrosis; acute respiratory distress syndrome;
                                                                                                                                                                                                                             Oligonucleotide associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airwa obstruction. The present sequence represents an oligonucleotide of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCR1, RANTES,
disease e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shahabuddin
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pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
     Homo sapiens.
                                                                                       pulmonary
                                                                                                                                                                                                                                                                                       06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                   ADJ59764 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CTCCCCAGTAGCTGGGATTA 20
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in S,
                                                                                       nypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 4 A;
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                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MCP4, useful for prophylaxis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ĕ,
                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sandrasagra A, H, Cong H;
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                                                                                    lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                             to RANTES
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Pred. No. 1.
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                                                                                                                                                                                                                                #13.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Miller
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 2; SEQ ID NO 620; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-203534/19.
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Shahabuddin S,
                                                                                                                                                                                                                                                                        obstruction.
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                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present invention
                                               537
                                                                                                18;
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                                                                                                                        Similarity
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CCTGCCTTAGCCTCCCGAGT
                                                 CCTGCCTCAGCCTCCCAAGT 556
                                                                                                Conservative
                                                                                                                                                                                                 BP; 2
                                                                                                                                                                                                                                                                        The present sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      asthma.
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4 F,
                                                                                                                                                                                                 A; 9 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        coding
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H, Cong H;
                                                                                                                     1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     relates to an oligonucleotide anti-sense to ing region with 2-10 nucleotides of 5'-end are
                                                                                                0
                                                                                                                     Score 16.8;
Pred. No. 1.
                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Aguilar
                                                                                                                        1.6e+03
                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ġ
                                                                                                                                                                                                                                                                             an
                                                                                                2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Miller
                                                                                                                                                                                                                                                                           oligonucleotide
                                                                                                                                             Length 20;
                                                                                                   Indels
                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           to e
                                                                                              Gaps
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interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; Oligonucleotide 06-MAY-2004 ADJ60955 standard; (first entry) associated DNA; 20 to PDE4C #21. 먪

05-FEB-2004. WO2004011613-A2 Homo sapiens pulmonary hypertension;

inflammation; bronchitis; oligonucleotide;

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RESULT 1452
ADJ60984
ID ADJ6098
XX ADJ6098
XC ADJ6098
XX Oligon
XX Oligon
XX interl
XX interl
XX interl
XX interl
XX eystic
XX gystic
XX W 2000
XX W0200
XX W0200
XX PN W0200
XX PP 05-FE
XX Y
PF 25-JU
XX EPIG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the coligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nyce JW, Ta
Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-JUL-2003; 2003WO-US023509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-203534/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JUL-2002; 2002US-0399076P
                                                                                                                                                                                                            interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; olig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (EPIG-)
                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                Oligonucleotide associated to PDE4C #50.
                                                                                                                                                                                                                                                                                                                                06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                              ADJ60984
                                                                                                    05-FEB-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nitiation codons and
                                    29-JUL-2002; 2002US-0399076P
                                                                    25-JUL-2003; 2003WO-US023509
        (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          791 GGGGTTCACCATGTTCGCCA 810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RANTES, MCP4, useful for prophylaxis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l Similarity
18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1
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                                                                                                                                                                                                                                                                                                                                                                                                standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGGTTTCACCATGTTGGCCA 20
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in S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lu
Lu
                                                                                                                                                                                                                                                                                                                                (first entry)
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H, Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                   inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.8; DB 1
Pred. No. 1.6e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Aguilar D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     or treating respiratory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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Sequence 20

BP; 4

Α.

6 C; 3

G; 7 T; 0 U;

0 Other;

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cc initiation codon, codding region with 2-10 nucleotides of 5 '-end and 3'-
cc end of nucleic acid target comprising gene(s) chosen from e.g.
cc interleukin (IL)-4 receptor, IL-5 receptor or salts of the
cc oligonucleotide and optionally surfactant operatively linked to the
cc oligonucleotide. The method is useful for preventing or treating a
cc respiratory or lung disease, which involves administering to the airways
cc of a subject an effective amount of an inhibitor. The oligonucleotide is
cc useful for production of a medicament for the prevention and/or treatment
cc of a respiratory or lung disease. The respiratory or lung disease is
cc chosen from airway inflammation, allergy(iss), asthma, impeded
cc respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases
cc (CODP), allergic rhinitis (AR), acute respiratory distress syndrome
cc (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway
construction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nyce JW, Ta
Shahabuddin
                                                                                                                                                                                                                                                                                                                                          The present invention relates to an oligonucleotide anti-sense to initiation codon, coding region with 2-10 nucleotides of 5'-end a
                                                                                                                                                                                                                                                                                                                                                                                                  Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                            disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  initiation
                                                                                                                                                                                                                                                                                                                                                                                                                                                         single or multiple target oligonucleotide anti-sense to e.g. ation codons and introns of respiratory disease-relevant genes e. RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 1840; 85pp;
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din S,
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Lu
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H, Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                  English.
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RESULT 1453
                                                                                                           Matches
                                                                                                                   Best Local
                                                                                                                         Query Match
06-MAY-2004
                ADJ59766;
                                ADJ59766 standard; DNA;
                                                                                         751 CACCACGCCTAGCTAATTTT 770
                                                                          \vdash
                                                                                                                   Similarity
                                                                          CACCATGCCTGGCTAATTTT
                                                                                                           Conservative
(first entry)
                                                                                                                  1.7%;
                                 20
                                 ΒP
                                                                                                           0
                                                                                                                Score 16.8; Ub 1,
                                                                           20
                                                                                                           Mismatches
                                                                                                                           DB 1;
                                                                                                           2
                                                                                                                          Length
                                                                                                           Indels
                                                                                                                            20;
                                                                                                           0,
                                                                                                           Gaps
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interleukin; IL-4 receptor; IL-5 receptor; lung disease;
airway inflammation; allergy; asthma; impeded respiration;
cystic fibrosis; acute respiratory distress syndrome;
pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
        Nyce JW, Tang
Shahabuddin S,
                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotide
                                                                                                          29-JUL-2002; 2002US-0399076P
                                                                                                                                                   25-JUL-2003; 2003WO-US023509
                                                                                                                                                                                           05-FEB-2004.
                                                                                                                                                                                                                                  WO2004011613-A2
                                                                                                                                                                                                                                                                          Homo sapiens
                                                                    (EPIG-) EPIGENESIS PHARM INC
Lu,
                                                                                                                                                                                                                                                                                                                                                                                                                                             associated to RANTES #15.
        Sandrasagra
H, Cong H;
                             Ą
                             Aguilar
                                Ď,
                                    Miller
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WPI; 2004-203534/19

Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.

e.g.,

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PROCESS OF A COLUMN AND A COLUM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1454
ADJ60973
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases
disease e.g.,
                   Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes (CR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                               05-FEB-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cystic fibrosis; acu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligonucleotide associated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to an oligonucleotide anti-sense to e. initiation codon, coding region with 2-10 nucleotides of 5'-end and end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (II)-4 receptor, II-5 receptor or salts of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim
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                                                                                                                                                                                                                                                                                                                                                                        25-JUL-2003; 2003WO-US023509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          interleukin; IL-4 receptor; IL-5 receptor; lung disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADJ60973;
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                                                                                                                                                                                                                                                                     (EPIG-)
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                                                                                                                                                                                                                                                                     EPIGENESIS
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                                                                                                                                                                                  Tang
in S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                     Tu,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             acute respiratory distresension; lung inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                  Sandrasagra
H, Cong H;
                                                                                                                                                                                                                                                                     PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 allergy; asthma; impeded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           90.0%;
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          distress syndrome;
mmation; bronchitis; oligonucleotide;
                                                                                                                                                                                                           Aguilar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    #39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .6e+03
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                                                                                                                                                                                                              'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    respiration;
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Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.

e.g.,

WPI; 2004-203534/19

Shahabuddin мусе JW,

Tang lin S,

L, Sandrasagra Lu H, Cong H;

P

Aguilar

'n

Miller

initiation codon,

present invention relates to

coding region with

2-10

an oligonucleotide

nucleotide anti-sense t

ទ

and

Example 5;

SEQ ID NO 2512; 85pp; English.

25-JUL-2003; 2003WO-US023509

WO2004011613-A2.

29-JUL-2002; 2002US-0399076P

(EPIG-) EPIGENESIS PHARM INC.

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\begin{array}{c} \mathsf{CCC} \times \mathsf{S} \times \mathsf{PP} \\ \mathsf{PP} \\
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome;
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end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is

The present invention

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The present invention relates to an oligonuclectide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(e) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonuclectide and optionally surfactant operatively linked to the oligonuclectide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonuclectide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome
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Pred. No. 1
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ADJ60949
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Best Local
                                                                           The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (II)-4 receptor, II-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(les), asthma, impeded respiration, cystic fibrosis (CR), chronic obstructive pulmonary diseases (CODD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway that the control of the control 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                airway inflammation; allergy; abumma, _____cystic fibrosis; acute respiratory distress syndrome; cystic fibrosis; acute respiratory distress syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes (CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Shahabuddin
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.6e+03;
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Query Match Best Local Matches

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cc initiation codon, coding region with 2-10 nucleotide anti-sense to e.g., cc end of nucleic acid target comprising gene(s) chosen from e.g. cc interleukin (IL)-4 receptor, IL-5 receptor or salts of the cc oligonucleotide and optionally surfactant operatively linked to the cc oligonucleotide and optionally surfactant operatively linked to the cc origonucleotide. The method is useful for preventing or treating a cc respiratory or lung disease, which involves administering to the airways cc of a subject an effective amount of an inhibitor. The oligonucleotide is cc useful for production of a medicament for the prevention and/or treatment cc fa respiratory or lung disease. The respiratory or lung disease is cc chosen from airway inflammation, allergy(ies), asthma, impeded cc respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (CC (COPD), allergic rhinitis (AR), acute respiratory distress syndrome cc (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway co bstruction. The present sequence represents an oligonucleotide of the cc invention.
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ADJ60972
ID ADJ6097
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Best Local Similarity
Matches 18; Conser
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                                                                                                                                                                                                                                                                                                                    Claim 2; SEQ ID NO 1828; 85pp; English
                                                                                                                                                                                                                                                                                                                                                                Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nyce JW, Tang L, Sandrasagra Shahabuddin S, Lu H, Cong H;
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Pred. No. 1.6e+03;
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ADJ59202
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                                                                                                                                                 The present invention relates to an oligonucleotide anti-sense to e.g., CC initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-CC end of nucleic acid target comprising gene(s) chosen from e.g. CC interleukin (II)-4 receptor, II-5 receptor or salts of the CC oligonucleotide and optionally surfactant operatively linked to the CC oligonucleotide. The method is useful for preventing or treating a CC respiratory or lung disease, which involves administering to the alivays CC of a subject an effective amount of an inhibitor. The oligonucleotide is CC useful for production of a medicament for the prevention and/or treatment CC of a respiratory or lung disease. The respiratory or lung disease is CC chosen from airway inflammation, allergy(ies), asthma, impeded CC respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (CODD), allergic rhinitis (NR), acute respiratory distrees syndrome CC (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway construction. The present sequence represents an oligonucleotide of the
                                                                     Query Match
Best Local :
                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        airway inflammation; allergy; asthma; impeded respiration;
cystic fibrosis; acute respiratory distress syndrome;
pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nyce JW, Ta
Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide associated to IL 4R
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                                                                                                                                                                                                                                                                                                                                                                                         Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-JUL-2002; 2002US-0399076P
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                                                                                                               Sequence
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                                                                     Similarity
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                                                                                                                                                                                                                                                                                                                                                                                         SEQ
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in S,
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                GCTGGGATTACAGGCGTGAG 883
                                                                                                               20
 GCTGGGATTATAGGCATGAG
                                                                                                               BP; 5
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                                                                                                                                                                                                                                                                                                                                                                                         ID NO 58; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 . m.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    receptor; IL-5 receptor; lung disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sandrasagra
H, Cong H;
                                                                   1.7%;
                                                                                                               2 C; 8
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                                                                                                               G; 5
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                                                       Score 16.8; Di
Pred. No. 1.6e
O; Mismatches
 20
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                                                                      .6e+03
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                                                          Gaps
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밁 S

542 CTCAGCCTCCCAAGTAGCTG 561

0

1 CTTAGCCTCCCGAGTAGCTG

20

RESULT 1462 ADJ59770

ADJ59770 standard; DNA;

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ВP

ADJ59770

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ADJ59765

ID ADJ59765 standard; D

XX

AC ADJ59765;

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DT 06-MAY-2004 (first

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DT 06-MAY-2004 (first

XX

Coligonuclectide asso

XX

Interleukin; IL-4 re

KW airway inflammation;

KW cystic fibrosis; acu

KW cystic fibrosis; 2003WO-

XX

Cys 29-JUL-2002; 2002US-

XX

Cys 29-JUL-2002; 2002US-

XX

Cys JW, Tang L, S

PI Nyce JW, Tang L, S

PI Shahabuddin S, Lu H

XX

Nyce JW, Tang L, S

PI Nyce
                                                                                                                                                                                                                     The present invention relates to an oligonucleotide anti-sense to e.g., cc initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g.

Cc interleukin (IL)-4 receptor, IL-5 receptor or salts of the city of the coligonucleotide and optionally surfactant operatively linked to the coligonucleotide. The method is useful for preventing or treating a cc respiratory or lung disease, which involves administering to the airways cf a subject an effective amount of an inhibitor. The oligonucleotide is cc useful for production of a medicament for the prevention and/or treatment cc useful for production of a medicament for the prevention and/or treatment cc arespiratory or lung disease. The respiratory or lung disease is cc chosen from airway inflammation, allergy(ies), asthma, impeded cc respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (CODD), allergic rhinitis (AR), acute respiratory distress syndrome cc (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway construction. The present sequence represents an oligonucleotide of the invertice.
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   Matches
                                   Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nyce JW, Tang L,
Shahabuddin S, Lu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide associated to RANTES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-203534/19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; SEQ ID NO 621; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (EPIG-) EPIGENESIS PHARM INC
18;
                                      Similarity
   Conservative
                                                                                                                                     BP; 3 A; 7 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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H, Cong H;
                                   1.7%;
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       0;
                                          Score 16.8; DB 1
Pred. No. 1.6e+03
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          Mismatches
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                                                                         DB 1;
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          2
                                                                         Length
          Indels
                                                                             20;
          0;
          Gaps
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ADJ60943
ADJ60943
ID ADJ6094
XX ADJ6094
AC ADJ6094
XX 06-MAY-
XX Oligonu
XX
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                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 18
                   Oligonucleotide associated to PDE4C #9.
                                                   06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
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                                                                                                                   ADJ60943 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                      Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; SEQ ID NO 626; 85pp; English.
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                                                                                                                                                                                                                                                                   l Similarity
18; Conserv
                                                                                                                                                                                                                                    GCAGTGGCGCAATCTTGGCT 675
                                                                                                                                                                                                     GCAGTGGCGCGATCTCGGCT 20
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in S,
                                                                                                                                                                                                                                                                                                                                   BP; 2 A;
                                                                                                                                                                                                                                                                     Conservative
                                                  (first entry)
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H, Cong H;
                                                                                                                                                                                                                                                                                                                                 6 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                    90.0%;
                                                                                                                                                                                                                                                                                                      1.7%;
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                                                                                                                                                                                                                                                                                   Score 16.8; DB 1
Pred. No. 1.6e+03
                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                      DB 1;
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                                                                                                                                                                                                                                                                                                  Length 20;
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RESULT 1464
ADJ59869
ID ADJ5986
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AC ADJ5986
XC ADJ5986
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O6-MAY-
DT O6-MAY-
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DE Oligonu
XX
interle
XW interle
XW airway
XW cystic
XW pulmona
XW bs:
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Best Local S
Matches 18
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                             airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                               Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 4 A; 10 C; 2 G; 4 T; 0 U; 0 Other;
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Shahabuddin S,
                                                                                                                          interleukin; IL-4 receptor; IL-5 receptor; lung disease;
                                                                                                                                                                                                                                                              06-MAY-2004
                                                                                                                                                                                                                                                                                                                             ADJ59869;
                                                                                                                                                                                                                                                                                                                                                                                         ADJ59869 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disease e.g., asthma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity
18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GCTCACTGCAACCTCTGCCT 692
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GCTCACTGCAACCTCCACCT
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                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                         DNA;
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H, Cong H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.8; DB 1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         represents an oligonucleotide of the
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RESULT 1465
ADJ60993
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonuclectide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to an oligonucleotide anti-sense to eninitiation codon, coding region with 2-10 nucleotides of 5'-end and end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a consistence of the method is useful for preventing or treating a consistence of the method is useful for preventing or treating a consistence of the method is useful for preventing or treating a consistence of the method is useful for preventing or treating a consistence of the method is useful for preventing or treating a consistence of the method is useful for preventing or treating a consistence of the method is useful for preventing or treating a consistence of the c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCRI, RANTES, MCP4, useful for prophylaxis or treating respiratory
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                                                                                                                                                                                 interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 2; SEQ ID NO 725; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-203534/19
                                                                                                                                                             pulmonary
                                                                                                                                                                                                                                                                                              Oligonucleotide associated to PDE4C #59.
                                                                                                                                                                                                                                                                                                                                                    06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADJ60993 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              obstruction.
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                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  647 GGCTGGAGTGCAGTGGCGCA 666
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18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              allergic rhinitis (AR), acute respiratory distress syndrome pulmonary hypertension, lung inflammation, bronchitis, airway tion. The present sequence represents an oligonucleotide of the
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lin S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GGCTGGAGTGAAGTGGCACA 20
                                                                                                                                                             hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                 (first entry)
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Lu H, Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.7%;
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                                                                                                                                                                lung
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>,</u>
                                                                                                                                                                inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16.8;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
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                                                                                                                                                                                                                                                                                               interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCR1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nyce JW, Ta
Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-JUL-2003; 2003WO-US023509
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 2; SEQ ID NO 1849; 85pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide
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                                                                                                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18;
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in S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                associated to PDE4C #60.
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Pred. No. 1.
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RESULT 1467
ADK41378/c
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the inventor of the control of the co
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                                                                       27-JUN-2002; 2002DK-00001005
07-OCT-2002; 2002DK-0000150
25-FEB-2003; 2003DK-00000289
29-APR-2003; 2003DK-00000639
                                                                                                                                                                                                                                                                                                                                                                                                               Human chromosome 19 RAI il
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADK41378 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (EPIG-)
              (UYAA-)
(ARBE-)
                                                                                                                                                                                                                08-JAN-2004.
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                                                                                                                                                                       27-JUN-2003; 2003WO-DK000448
                                                                                                                                                                                                                                                       WO2004003229-A2
                                                                                                                                                                                                                                                                                                                                                   eingle
                                                                                                                                                                                                                                                                                                                                                                        sequence polymorphism analysis;
                                                                                                                                                                                                                                                                                                                sapiens
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                                                                                                                                                                                                                                                                                                                                                     nucleotide
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              UNIV AARHUS.
ARBEJDSMILJO
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in S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
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                                                                                                                                                                                                                                                                                                                                                   polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
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H, Cong H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
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                INST NAT
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                                                                                                                                                                                                                                                                                                                                                                                                               sensor probe.
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Pred. No. 1.
                  INST OCCUPA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                     human; chromosome SNP; probe.
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                                                                                                                                                                                                                                                                                                                                                                          cancer;
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RESULT 1468
ADK41252/c
ID ADK4125
XX
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer to a disease treatment; a primer or probe for use in the method of estimating the disease risk or prognosis of an individual or for estimating a treatment response of an individual suffering from cancer to a disease treatment; an antibody directed to an epitope of a RAI gene product; and a kit for use in the method of estimating the disease risk or prognosis of an individual or for estimating a treatment response of an individual suffering from cancer to a disease treatment, comprising at least one primer or probe and optionally amplifying means for nucleic acid amplification. The novel method is useful for estimating the disease risk or prognosis of an individual or for estimating a treatment response of an individual suffering from cancer to a disease treatment. This polynucleotide sequence represents a probe used in the exemplification of the invention.
                                                                                                                                                                                                                                                       27-JUN-2002;
07-OCT-2002;
25-FEB-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence polymorphism analysis;
single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel method of estimating disease risprognosis of an individual by sequence polymorphism analysis, esp polymorphisms in the human chromosome 19q. The invention further to: estimating a treatment response of an individual suffering from the content of the content 
Estimating the disease polymorphism analysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 136; 145pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Estimating the disease polymorphism analysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nexo
                                                                      WPI; 2004-142878/14
                                                                                                                    Nexo
                                                                                                                                                                                                                                                                                                                                                        27-JUN-2003; 2003WO-DK000448
                                                                                                                                                                                                                                                                                                                                                                                                   08-JAN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADK41252;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADK41252
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2004003229-A2
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                                                                                                                    BA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       480
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18; Conserv
                                                                                                                                                                UNIV AARHUS.
ARBEJDSMILJO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GTGCAGTGGTGTGATCACAG 499
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                                                                                                                 Vogel U,
                                                                                                                                                                                                                                   2002DK-00001005.
2002DK-00001500.
2003DK-00000289.
2003DK-00000639.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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                                                                                                                      Rockenbauer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                    INST
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ۲
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                                                                                                                    'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human; chromosome 19q; SNP; probe; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ĮIJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 English.
                                                                                                                                                                    OCCUPA.
                                                                                                                      Bukowy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .6e+03;
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                                                                                                                    ZK
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                             individual by sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        risk or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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risk

or.

prognosis

of an

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of breast cancer-1 such as a hyperproliferative disorder in particular breast, ovary, prostate and peritoneum cancers. The invention is also used in antisense therapy. The present sequence is human breast cancer associated antisense oligonucleotide. Note: This sequence given in the sequence listing differs from that given in example 15 of the

cancer-1

sequence listing differs from that given in

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RESULT 1469
ADJ96296
ID ADJ9629
XX ADJ9629
XX ADJ9629
XX Human b
XX Unident
XX Unident
XX Unident
XX US2004(
XX US2004(
XX Ha-JUL-
XX IB-JUL-
XX IB-JUL-
XX IB-JUL-
XX WPI; 21
XX Unident
XX US2004(
XX IB-JUL-
X
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Claim 30; SEQ IJ NO 10; 145pp; English

The invention relates to a novel method of estimating disease risk or cc prognosis of an individual by sequence polymorphism analysis, especially colymorphisms in the human chromosome 19q. The invention further relates to: estimating a treatment response of an individual suffering from cc cancer to a disease treatment; a primer or probe for use in the method of catinating a treatment response of an individual suffering from cc estimating a treatment response of an individual suffering from cancer to a disease treatment; an antibody directed to an epitope of a RAI gene cc product; and a kit for use in the method of estimating the disease risk or prognosis of an individual or for estimating a treatment response of can individual suffering from cancer to a disease treatment, comprising at least one primer or probe and optionally amplifying means for nucleic caid amplification. The novel method is useful for estimating the disease cf isk or prognosis of an individual or for estimating a treatment response of constant of promocancer to a disease treatment response considered to a nidividual suffering from cancer to a disease treatment response considered to a nidividual suffering from cancer to a disease treatment response considered to a nidividual suffering from cancer to a disease treatment of the considered to a nucleotide sequence represents a primer/probe used for detecting the disease treatment of invention.

Sequence 20 BP; 5 A, 8 C; 4 G; 3 T; 0 U; 0 Other;

Query Match Best Local S Matches 18 18; Similarity Conservative 1.7%; <u>.</u>. Score 16.8; Pred. No. 1 Mismatches ..6e+03; DB 1; Length Indels 0 Gaps 0

ADJ96296 standard; DNA;

20

06-MAY-2004 (first entry)

Human breast cancer-1 associated antisense oligonucleotide

antisense therapy; diagnosis; hyperproliferative antisense; disorder;

Synthetic. Unidentified

US2004014051-A1.

22-JAN-2004.

18-JUL-2002; 2002US-00199676

18-JUL-2002; 2002US-00199676

SISI PHARM

Brown-Driver VL, Dobie 즟

WPI; 2004-121557/12.

preventing and/or treating
cancer-1, such as breast, New antisense oligonucleotide compounds, useful for diagnosing, d/or treating conditions with aberrant activity of breast h as breast, ovary, prostate and/or peritoneum cancers.

Disclosure; SEQ ID NO 37; 175pp; English.

The present invention is directed to novel antisense compounds targett to breast cancer-l proteins and their encoding nucleic acids. The invention is useful for the diagnosis, prevention and/or treatment of diseases and conditions associated with aberrant expression and activity expression and activity targetted

Ś

1058 ACACCCCGCTAATTTTTGTA 1077

Query Match Best Local S Matches 18

Similarity

1.7%;

Score 16.8; Pred. No. 1.

.6e+03

DB

1:

Length

20;

0

Mismatches

2

Indels

0;

Gaps

0

RESULT 1470
ADJ96332/c
ID ADJ9633
XX ADJ9633
XX DT 06-MAYXX Breast
KW Breast
KW Breast
KW Breast
KW Breast
KW Breast
KW Breast
XX US20040
PN US20040
XX US20040
XX US20040
XX HB-JULXX ន្តម្ភិន្តិនិន្តនិង 멼 S Query Match Best Local S Matches of breast cancer-1 such as a hyperproliferative disorder in particular breast, ovary, prostate and peritoneum cancers. The invention is also used in antisense therapy. The present sequence is human breast cancer-associated antisense oligonucleotide. Note: This sequence given in the sequence listing differs from that given in example 15 of the New antisense oligonucleotide compounds, useful for diagnosing, preventing and/or treating conditions with aberrant activity of cancer-1, such as breast, ovary, prostate and/or peritoneum can Synthetic. Unidentified Sequence 20 BP; 4 A; 5 C; 4 G; 7 T; 0 U; 0 Other; to breast cancer-1 proteins and their encoding nucleic acids. The invention is useful for the diagnosis, prevention and/or treatment of diseases and conditions associated with aberrant expression and activity Breast cancer-1; diagnosis; hyperproliferative disorder; cancer; 06-MAY-2004 ADJ96332; ADJ96332 standard; WPI; 2004-121557/12. Brown-Driver VL, Dobie 18-JUL-2002; 2002US-00199676 18-JUL-2002; 2002US-00199676 22-JAN-2004. antisense therapy; antisense; Sequence 20 sequence listing differs specification. The present invention is directed to novel antisense compounds Disclosure; SEQ ID NO 73; 175pp; English. US2004014051-A1 1058 ACACCCCGCTAATTTTTGTA 1077 breast 18; μ SISI Similarity ACGCCCGGCTAATTTTTGTA BP; Conservative cancer-1 associated PHARM INC. (first entry) 7 A; DNA; 4 C; 90.0%; 20 տ ВP <u>,</u> G; Score Pred. 88 20 4 Mismatches T; antisense oligonucleotide No. 16.8; 0 ď; .6e+03 0 DB Other; 1. <u>ې</u> Length 20; Indels 0, cancers. targetted Gaps 0

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ADJ96456/c
ID ADJ964
XX ADJ964
XX ADJ964
XX Breast
XX ISIS-
XX I
RESULT 1472
ADJ96392
ID ADJ9639
XX
AC ADJ9639
XY
DT 06-MAY-
XX
DE Human L
XX
Breast
KW Breast
KW antiser
XX
Homo 86
OS Synthet
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                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention is directed to novel antisense compounds targetted to breast cancer-1 proteins and their encoding nucleic acids. The invention is useful for the diagnosis, prevention and/or treatment of diseases and conditions associated with aberrant expression and activity of breast cancer-1 such as a hyperproliferative disorder in particular breast, ovary, prostate and peritoneum cancers. The invention is also used in antisense therapy. The present sequence is human breast cancer-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotide compounds, useful for diagnosing, preventing and/or treating conditions with aberrant activity of breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Brown-Driver VL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Breast cancer-1; diagnosis; hyperproliferative disorder; cancer; antisense therapy; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human breast cancer-1 target oligonucleotide #41.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADJ96456 standard; DNA; 20
    Homo sapiens
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 7 A; 4 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-JUL-2002; 2002US-00199676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-JUL-2002; 2002US-00199676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US2004014051-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 15; Page 32; 175pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-JAN-2004
                                                                                                                        Human breast cancer-1 antisense oligonucleotide #197041
                                                                                                                                                                                                       ADJ96392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ISIS-) ISIS PHARM INC.
                                                              antisense therapy;
                                                                                                                                                                                                                                               ADJ96392 standard;
                                                                                                                                                                                                                                                                                                                                                                       1058 ACACCCCGCTAATTTTTGTA 1077
                                                                                                                                                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20
                                                                                                                                                                                                                                                                                                                                                                                                                               18;
                                                                              cancer-1; diagnosis; hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                ACGCCCGGCTAATTTTTGTA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACGCCCGGCTAATTTTTGTA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    such as breast, ovary, prostate and/or peritoneum cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dobie
                                                                                                                                                                                                                                               DNA;
                                                                human; antisense;
                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ₹,
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                                                                                                                                                                                                                                               BP.
                                                                                                                                                                                                                                                                                                                                                                                                                             <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 16.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  .6e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                               2
                                                                                   disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                   cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                             ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                               0,
        AX FX BX BX BX BX BX
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RESULT 1473
ADL14967
                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          to breast cancer-1 proteins and their encoding nucleic acids. The invention is useful for the diagnosis, prevention and/or treatment of diseases and conditions associated with abscrant expression and activity of breast cancer-1 such as a hyperproliferative disorder in particular breast, ovary, prostate and peritoneum cancers. The invention is also used in antisense therapy. The present sequence is human breast cancer-1 antisense oligonucleotide. Note: This sequence given in example 15 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonuclectide compounds, useful for diagnosing, preventing and/or treating conditions with aberrant activity of breast cancer-1, such as breast, ovary, prostate and/or peritoneum cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Brown-Driver VL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified_base
EP1388590-A2
                                     Homo sapiens
                                                                                                                 Human glaucoma-related optineurin
                                                                                                                                                         06-MAY-2004
                                                                                                                                                                                                ADL14967;
                                                                                                                                                                                                                                   ADL14967
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          specification differs from that given
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention is directed to novel antisense compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 15; Page 31; 175pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-JUL-2002; 2002US-00199676.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-JUL-2002; 2002US-00199676.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-JAN-2004.
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                                                                                                                                                                                                                                                                                                                                                                1058 ACACCCCGCTAATTTTTGTA 1077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2004-121557/12
                                                                                                                                                                                                                                                                                                                                                                                                                        18;
                                                                                                                                                                                                                                                                                                                                         1 ACGCCCGGCTAATTTTTGTA 20
                                                                          glaucoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 BP; 4 A; 5 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                        standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mod_base= 0
/note= "2'- 1
16. .20
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/mod_base= 0
/note= "2'-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag=
/mod_ba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "mod_base= OTHER
note= "Phosphorothioate backbone where all cytidines
'- methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dobie KW,
                                                                          optineurin;
                                                                                                                                                                                                                                                                                                                                                                                                                                        1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ۵
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                                                                                                                                                                                                                                          ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                      Score 16.8; D
Pred. No. 1.6e
0; Mismatches
                                                                            OPTN;
                                                                                                                    (OPIN) exon
                                                                            diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          in the sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                          1.6e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                 6 PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                            primer; ss
                                                                                                                    SF6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     targetted
                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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RESULT 1474
ADL23335/c
ID ADL2333
XX
AC ADL2333
XX
DT 20-MAY-
XX
DT 20-AUG-
PR 26-AUG-
PR 26
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence is that of PCR primer SF6 for exon 6 ADL14952 of the glaucoma-associated gene, OPTN (optineurin) ADL14949. The invention relates to a gene assay method for predicting future onset of primary open angle glaucoma and/or normal ocular tension glaucoma. This involves detecting a mutation in the OPTN gene coding sequence, specifically a substitution of G for A at position 619 and/or a substitution of A for G at position 898 of the OPTN coding sequence. The mutation(s) is detected using a nucleic acid amplification method using primers specific for the different exons of the coding sequence, including primers SF6 and SR6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gene assay for predicting future onset of glaucoma, particularly primary open angle glaucoma or normal ocular tension glaucoma, comprises detecting a mutation of at least one base of the optineurin gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        allelic deletion; PHIT; fragile histidine triad gene; PR; progesterone receptor; DLEC1; deleted in lung and oesophageal cancer 1; TRIM29; tripartite motif-containing 29; microsatellite; D3S1300; D3S1260; D11S35; D11S528.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-146134/15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-JUL-2003; 2003EP-00447201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 9; SEQ ID NO 19; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ss; primer; diagnosis; cervical allelic deletion; FHIT; fragile
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SYSM-)
     WPI; 2004-226867/21:
                                                                                                                                                                                                                                                                                            04-MAR-2004
                                                                                                                                                                                                                                                                                                                                                 WO2004018711-A2
                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Primer #1 for amplification of D3S1611.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADL23335
                                                       Ming-Qing
                                                                                                                                                             24-AUG-2002;
26-AUG-2002;
                                                                                                                                                                                                                                        20-AUG-2003; 2003WO-GB003637.
                                                                                                           (UNIO ) UNIV
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18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TGCACCACTACACCTGGCTA 593
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for exon 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TGTGCCACTACACCTGGCTA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Masago A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                             2002GB-00019890
2002US-0405717P
                                                                                                           COLLEGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7 C; 4 G; 5 T;
                                                                                                           LONDON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Takahata
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 16.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Н
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         intraepithelial neoplasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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Diagnosing cervical intraepithelial neoplasia comprising detecting an allelic deletion in genes selected from FHIT, PR, DLEC1- or TRIM 29 by comparing the FHIT, PR, DLEC1 and/or TRIM 29 polynucleotides or proteins present in the samples.
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Disclosure; SEQ ID NO 17; 56pp; English

This sequence represents a primer which was used in the method of the cinvention for diagnosing susceptibility to persistence or progression of cervical intraepithelial neoplasia (CIN) in an individual suffering from the disease. The method comprises detecting an allelic deletion in one or the disease. The method comprises detecting an allelic deletion in one or core genes selected from FHIT (fragile histidine triad gene), PR CC (progesterone receptor), DLEC1 (deleted in lung and oesophageal cancer 1) CC or TRIM29 (tripartite mottif-containing 29) by comparing the FHIT, PR, CC DLEC1 and/or TRIM29 polynucleotides or proteins present in the samples contrived from non-dyskaryotic and dyskaryotic samples, respectively. The CC derived from non-dyskaryotic and dyskaryotic samples, respectively. The CC of primers, where each pair of primers is suitable for amplifying a CC of primers, where each pair of primers is suitable for amplifying a microsatellite DNA marker selected from D3S1300, D3S1260, D11S35 or CC microsatellite DNA marker selected from D3S1300, D3S1260, D11S35 or CC deletion forms of a polynucleotide or protein selected from FHIT, PR, CC TRIM29 or DLEC1. The method is useful for diagnosing susceptibility to CC persistence or progression of cervical intraepithelial neoplasia in an CC individual suffering from the disease.

Sequence 20 BP; 6 A; 6 C; 4 G; 4 T; 0 U; 0 Other;

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                                                                                                                                               RESULT 1475
                                                                                                                                                                                Query Match
Best Local S
                                                                                                                                                                             Matches
                                                                                                                              ADL81396;
                                                                                                                                       ADL81396 standard; DNA;
                                                                                                                      20-MAY-2004
                                                                                                                                                                 384 CTCCCAAAGTGCTGGGATTA 403
                                                                                                                                                            20
                                                                                                                                                                             18;
                                                                                                                                                                                 Similarity
                                                                                                                                                            CTCTCGAAGTGCTGGGATTA 1
                                                                                                                                                                             Conservative
                                                                                                                      (first entry)
                                                                                                                                                                                 90.0%;
                                                                                                                                        20
                                                                                                                                        ВP
                                                                                                                                                                             <u>.</u>
                                                                                                                                                                                 Score 16.8;
Pred. No. 1
                                                                                                                                                                             Mismatches
                                                                                                                                                                                 DB 1;
                                                                                                                                                                                      Length
                                                                                                                                                                              Indels
                                                                                                                                                                              0
                                                                                                                                                                              Gaps
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asthma; bronchial hyperresponsiveness; obesity;

disease;

human; gene 216; ss; primer.

inflammatory bowel

Keith T,

Little RD,

Eerdewegh

P۷,

Dupuis

'n

Del Mastro

RG;

(EERD/) (KEIT/)

SIMO/) DMAS/)

DEL MASTRO R SIMON J. ALLEN K. EERDEWEGH DUPUIS J.

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PANDIT

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13-APR-1999; 99US-0129391P 13-APR-2000; 2000US-00548797 13-APR-2001; 2001US-00834597

99US-0129391P

KEITH T.

LITTLE R

Ö ש

19-APR-2002; 2002US-00126022

05-FEB-2004. US2004023215-A1 Homo sapiens Gene 216 polymorphism sequencing

primer #52

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ARESULT 1476
ADK74414
ID ADK7441
XX ADAK741

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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New
nucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, triggeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate
                                                                                                                                                                                                                                                                                              New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                bowel disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to an isolated nucleic acid molecule, or a set of nucleic acid molecules each given in the specification. The composition and methods are useful in diagnosing or treating asthma or bronchial hyperresponsiveness, and other diseases such as obesity or inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Simon
                                                                                                                                                         The present invention relates to an antisense compound targeted to nucleic acid molecule encoding Navl 3 where the accompound targeted to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-AUG-2003; 2003WO-US025465.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2004016754-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diabetic neuropathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nav1.3; Analgesic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chimeric phosphorothioate oligonucleotide to target Nav1.3 #1748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADK74414;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADK74414 standard;
                                                                                                                                                                                                                                                 Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-AUG-2002; 2002US-0403416P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   isolated nucleic acid molecules useful for diagnosing or treating hma or bronchial hyperresponsiveness, or other diseases such as sity or inflammatory bowel disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                          2004-203785/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     686 TCTGCCTCCCGGGTTCAAGT 705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ģ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20
                                                                                                                                                                                                                                              SEQ ID NO 1748;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TCTGCCTCCCAGATTCAAGT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Allen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 4 A; 7 C; 3 G; 6 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>,</u>~
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nootropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          arthritic pain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               208; 485pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20
                                                                                                                                                                                                                                           417pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Neuroprotective; post-herpetic neuralgia; pain; migraine headache;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16.8;
No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .6e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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RESULT 1477
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                                                                                        Query Match
Best Local S
Matches 18
                                                                                                                                                                                 headache; seizure disorder such as childhood seizure disorder, inc
but not limited to meonatal or infantile epilepsy; or ataxia. The
sequence represents a chimeric phosphorothioate oligonucleotide wi
2'MOE wings and a deoxy gap. Used during the antisense inhibition
human Navl.3 expression, the oligonucleotides are designed to targ
                                                                                                                                             Sequence
                                                                                                                                                                      different regions of the human Nav1.3 RNA.
                                                                426
                                       <u>ب</u>
                                                                                          18;
                                                                                                       Similarity
                                                             CTTTTTATTTTATTTTTTT
                                                                                                                                             20 BP; 0
                                                                                          Conservative
                                                                                                                                             A; 1 C; 0 G; 19 T; 0 U;
                                                                                                    1.7%;
                                                                                          <u>,</u>
                                                                                                       Score 16.8;
Pred. No. 1
                                                                 445
                                       20
                                                                                           Mismatches
                                                                                                       1.6e+03;
                                                                                                                     띪
                                                                                                                                             0 Other;
                                                                                                                   1;
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                                                                                                                  Length
                                                                                           Indels
                                                                                                                     20;
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                                                                                          Gaps
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human; medicine; signal transduction; glycoprotein; transcription; oligo-capping method; ss; PCR; primer.
                                                                                                                                             EP1396543-A2.
                                                                                                                                                                         Clone specific PCR primer to amplify human full length cDNA SeqID 4410.
                                                                                                                                     10-MAR-2004.
                                                                                                                                                                                          ADL32377;
                                                                                                                                                                                                  ADL32377
                                                                                                                                                                                                  standard;
                                                                                                                                                                                  (first
                                                                                                                                                                                                  DNA;
                                                                                                                                                                                  entry)
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11-JAN-2000; 02-MAY-2000; 08-JUL-1999; 07-JUL-2000; (REAS-) RES ASSOC BIOTECHNOLOGY 99JP-00194486. 2000JP-00118774. 2000JP-00183865. 2000BP-00114089. 2003EP-00025638

New oligonucleotide primers (830 cDNAs) useful Wakamatsu A, 2004-204755/20 Sugiyama T, Isogai T, ai T, Hayashi K, Nagai K, Kojima တ Ishii S, S, Otsuki Kawai T, Ko Н;

for

synthesizing

Nishikawa

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Example 18; SEQ ID NO 4410; 1340pp; English

human cDNAs.

This invention relates to a novel primers useful for synthesising full length cDNA molecules that encode human proteins. Specifically, it refers to secretory or membrane proteins that are potential therapeutic agents/ target molecules in the field of medicine, and in particular genes encoding proteins that are associated with signal transduction, glycoproteins and transcription. The present invention describes a method for efficiently cloning a full length human cDNA from both the 5° and 3° ends using the oligo-capping method. This oligonucleotide sequence is a human clone specific PCR primer used in an exemplification of the

Sequence 20 ₿P; ហ Þ 7 C; 4 G; 4 T; 0 U; 0 Other;

Query Match
Best Local Similarity
Matches 18; Conserv Conservative 90.0%; Score 16.8; D Pred. No. 1.6e 0; Mismatches 0 6e+03 DB 1; Length Indels 20 0, Gaps

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RESULT 1479
ADL34877/c
ID ADL3487
XX
AC ADL3487
XX
DT 17-JUN-
XX
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                                                                                                                                                     RESULT 1478
ADL32402
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                                                                                               ş
                                                                                                                  Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                          This invention relates to a novel primers useful for synthesising full length cDNA molecules that encode human proteins. Specifically, it refers to secretory or membrane proteins that are potential therapeutic agents/ target molecules in the field of medicine, and in particular genes encoding proteins that are associated with signal transduction, glycoproteins and transcription. The present invention describes a method for efficiently cloning a full length human cDNA from both the 5' and 3' ends using the oligo-capping method. This oligonucleotide sequence is a human clone specific PCR primer used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                   08-JUL-1999; 99JP-00194486.
11-JAN-2000; 2000JP-00118774.
02-MAY-2000; 2000JP-00183865.
07-JUL-2000; 2000EP-00114089.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADL32402 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Clone specific PCR primer to
                                                                                                                                                                                                                                                                                                                                                                                                             07-JUL-2000; 2003EP-00025638
                                                                                                                                                                                                                                                                                                                                                                                                                                               EP1396543-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human; medicine; signal transduction; glycoprotein; transcription;
oligo-capping method; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-MAY-2004
                                                                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                        New oligonucleotide primers (830 cDNAs) useful
                                                                                                                                                                                                                                                                                                                                  Ota T,
                                                                                                                                                                                                                                                                                                                                                                                                                              10-MAR-2004
        17-JUN-2004
                                                                                                                                                                                                                                                               Example 18; SEQ ID NO 4435; 1340pp; English
                                                                                                                                                                                                                                                                                                                         Wakamatsu A,
                                          ADL34877
                                                                                                                                                                                                                                                                                                                                                   (REAS-) RES ASSOC BIOTECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                               sapiens
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                                                                                                  542
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                                                                                                                                                                                                                                                                                 human cDNAs.
                                                                                                                                                                                                                                                                                                                                  Nishikawa T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTCAGCCTCCCAAGTAGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTCAGCCTTCCAAGTAGCAG
                                                                                              CTCAGCCTCCCAAGTAGCTG
                                                                                   CTCAGCCTTCCAAGTAGCAG 20
                                          standard;
                                                                                                                                                     BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                    Conservative
         (first entry)
                                                                                                                                                                                                                                                                                                                         Sugiyama
                                                                                                                                                     5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                          DNA;
                                                                                                                         1.7%;
                                                                                                                                                     7 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                  Isogai T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                           <u>გ</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                          ni T, Hayashi K,
Nagai K, Kojima
                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amplify human full length cDNA SeqID 4435
                                                                                                                            Score 16.8;
Pred. No. 1
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                                                                                                    561
                                                                                                                    Mismatches
                                                                                                                            1.6e+03;
                                                                                                                                     DB
                                                                                                                                                                                                                                                                                                                          Ishii S,
1 S, Otsuki
                                                                                                                                    1;
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                                                                                                                                  Length 20;
                                                                                                                     Indels
                                                                                                                                                                                                                                                                                         synthesizing
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T, Koga
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                                                                                                                    Gaps
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RESULT 1480
ADL34724
                                                                                                                                                                                                                                          Matches
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes novel antisense oligonucleotides targeted to a nucleic acid encoding PPAR-delta, which specifically hybridise to and inhibit expression of PPAR-delta. The oligonucleotide specifically hybridises with at least an 8-nucleobase portion of an active site on the nucleic acid molecule encoding the PPAR-delta and comprises at least one modified internucleoside linkage, which is a phosphorothioate linkage, at least one modified sugar moiety, which is a 2'-O-methoxyethyl sugar moiety or at least one modified nucleobase, which is a 5-methylcytosine. The antisense oligonucleotides are useful for preparing a composition for treating hyperproliferative disorders, e.g., cancer. The oligonucleotides of the invention have cytostatic activity and can be used for gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotide, having a sequence targeted to acid encoding PPAR-delta, useful for preparing a composition hyperproliferative disorder, e.g., cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antisense; PPAR-delta; human; hybridisation; inhibitor; phosphorothioate linkage; 2'-O-methoxyethyl sugar; 5-methylcytosine; hyperproliferative disorder; cancer; cytostatic; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human PPAR-delta target site ID 49985
                                                                                                                                                                                                                                                                                                                                             Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaarde
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GAAR/) GAARDE W. (FREI/) FREIER S (WATT/) WATT A T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-MAY-2002; 2002US-00160807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-SEP-2003; 2003US-00655847
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
ADL34724 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 16; SEQ ID NO 175; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-282460/26.
                                                                                                                                                                                      1027 CAAGCAGCTGGGATTACGGG 1046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E,
                                                                                                                                 20 CAAGTAGCTGGGATTACAGG 1
                                                                                                                                                                                                                                          18;
                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Freier SM,
                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                             BP; 4 A; 7 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3
                                                                                                                                                                                                                                                                      90.0%;
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  股
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AT.
                                                                                                                                                                                                                                          <u>,</u>
                                                                                                                                                                                                                                                                                               Score 16.8;
                                                                                                                                                                                                                                                                      Pred.
                                                                                                                                                                                                                                             Mismatches
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for treating
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antisense; PPAR-delta; human; hybridisation; inhibitor; phosphorothioate linkage; 2'-O-methoxyethyl sugar; 5-methylcytosine; hyperproliferative disorder; cancer; cytostatic; gene therapy; ss;

Antisense oligonucleotide ISIS

136865

17-JUN-2004 ADL34724;

(first entry)

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US2004063129-A1

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                                                                                                                                                                                                                                                                                                                                       immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological;
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                                                                                                                                                                                                                                                        Synthetic.
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(FREI/) FREIER S M.
(WATT/) WATT A T.
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/*tag= b
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/note= "phosphorothioate linkages
/maidues are 5-methylcytidines"
                                                                                                                                                                                           Location/Qualifiers
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Pred. No. 1.
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                               antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition associated with mPGES-1 e.g., isbhaemia or reperfusion injury, or disease, arthritis, diabetes, cancer, isbhaemia or reperfusion injury, or
chimeric; antisense oligonucleotide; phosphorothioate; human;
microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor;
microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                                 Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1224
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                                                                                                                                                                                                                                                                                  TGCTGGGATTACAGGCGTGA 882
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/mod_ba
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/note= "2'-O-methoxyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. Chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,
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                                                                                1.7%;
                                                                                                                                                                   6 C; 1
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                                                                                                                                                                      G; 5 T; 0 U; 0 Other;
                                                                                Score 16.8;
Pred. No. 1
                                                            Mismatches
                                                                                      1.6e+03
                                                                                                                DB 1;
                                                                                                             Length
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RESULT 1483
ADM15443/c
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                 The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cardiovascular
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                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a encoding mPGES-1, useful for preparing a composition for inflammation, Alzheimer's disease, arthritis, diabetes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-APR-2004.
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                                                                                                                                                                                                                                                                                                                       Claim 4; SEQ
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antidiabetic,
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                                                                                                                                                                                                                                                                                                                       ID NO 1630; 132pp; English.
  immunomodulator,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disorder;
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/mod_base= OTHER
/note= "2'-O-methocyethyls"
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/mod_base= OTHER
/mote= "2'-O-methoxyethyls"
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  cardiant,
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                                                                                                                                                                                                                                                                                                                                                                                                          es, cancer or
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological;
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New antisense compound, having a sequence targeted to a nucleic ace encoding mPGES-1, useful for preparing a composition for treating
                                                                                                                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                             modified_base
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                                           WPI; 2004-305094/28.
                                                                                                                                                                                                                    WO2004028458-A2
                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                               25-SEP-2002; 2002US-0413549P
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                                                                                                  (PHAA ) · PHARMACIA CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                 /*tag= a
/mod_base= OTHER
/note= "2'-O-methocyethyls"
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/note= "phosphorothioate linkages
residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                note=
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                                                                                                                                                                                                                                                              base= OTHER
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Pred.
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No. 1.
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 e.g.,
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9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                         can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury,
                                                                                                                                                                                                                                                                                                                                           human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compound
                                                                                                                            antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound
                                        ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                       targeted
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20 BP; 4
                                          immunological,
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  4 C;
    10
                                        cardiovascular or neurological disorder
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  N
    Ή.
  0 U;
    0 Other;
                                                                                                                                                                                                                                                                                                                                                                                       antisense oligonucleotide
E2 synthase (mPGES-1). Th
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Matches Query Match Best Local ADM14625 711 20 18; Similarity TCCTGCCCCAGCCTCCTGAG 730 standard; DNA; TCCCGCCTCAGCCTCCTGAG 1 Conservative 1.7%; 20 ВP 0 Score 16.8; Pred. No. 1 Mismatches 6e+03 Indels 0, Gaps

DB 1;

Length

0,

or R

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01-JUL-2004 ADM14625;

(first entry)

chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase; mpGES-1 inhibitor; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorcardiovascular disorder; neurological disorder; ss. Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:812.

Homo sapiens Synthetic.

ADM14625/c
ID ADM14625/c
ID ADM1462
XX ADM1462
XX ADM1462
XX ADM1462
XX Chimeri
KW chimeri
KW microso
KW micro modified\_base modified\_base modified\_base residues /\*tag= /mod\_base= OTHEk /note= "2'-O-methocyethyls" /note= "phosphorothioate linkages
residues are 5-methylcytidines" Location/Qualifiers 16. .20 mod\_base= OTHER \*tag= b \_base= OTHER e= "2'-O-methoxyethyls" ი ρ and all cytidine

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ADM14799/c
ID ADM147
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Matches
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                                                   chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antiinflammatory, neuroprotective, nootropic, antiinflammatory, neuroprotective, nootropic, antiinflammatory, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targetted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric having a disease or condition associated with mPGES-1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20
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                                                                                                                                                                                                                                               Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:986
                                                                                                                                                                                                                                                                                                                          ADM14799
                                                                                                                                                                                                                                                                                                                                                              ADM14799 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.8;
Pred. No. 1
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XSX

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                                                                                                                                                                                                                                                                 antisense oligonuclectides and antisense compounds have cytostatic, antidisabetic, immunomodulator, cardiant, neuroprotective, antidiflammatory, neuroprotective, nootropic, antiarthirtic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                     676 CACTGCAACCTCTGCCTCCC 695
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                                                                                                                                     Similarity
CACTGCAGCCTCCGCCTCCC 1
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                                                                                                        Conservative
                                                                                                                                                                                                                   3 A;
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residues are 5-methylcytidines"
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/note= "2'-O-methoxyethyls"
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                                                                                                                                                                                                                   3 C; 12 G; 2 T; 0 U; 0 Other;
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                                                                                                                                     Score 16.8;
Pred. No. 1
                                                                                                             Mismatches
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01-JUL-2004

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The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antidiabetic, immunomodulator, cardiant, neuroprotective, candiabetic, immunomodulator, and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mpGES-1 residence compound CC can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC conthibution associated with mpGES-1 e.g., instammation, Alzheimer's compound controlics artirities, diabetes, cancer, ischaemia or reperfusion injury, or on the lamic remained controlics and cardiovascular activities, and can be used for preparing a composition for treating a disease or conthibutor and in gene therapy. The antisense compound CC content activities, diabetes, cancer, ischaemia or reperfusion injury, or on the propertusion injury, or content activities and cardiovascular activities.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human mPGES-1 chimeric antisense oligonucleotide
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/note= "phosphorothioate linkages and all
residues are 5-methylcytidines"
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/note= "2'-O-methocyethyls"
16. .20
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        cardiovascular or
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neurological
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Sequence 20

BP; 4 A; 8 C;

4 G; 4 T; 0 U; 0 Other;

describes:

antisense

compounds

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human mi
9q34.3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1 inpEES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological;
                                                                                                                                                           New antisense compound, having a sequence targeted to a nucleic avenceding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                     The present sequence represents a chimeric targeted to human microsomal prostaglandin
                                                                                                                                                                                                                                                WPI; 2004-305094/28
                                                                                                                                                                                                                                                                                      Gierse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
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                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                      25-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2004028458-A2
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                                                                                                                                                                                                                                                                                                                              (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mmunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sapiens.
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mPGES-1 gene is located on chromosome 3. The present invention also describes
                                                                                                     4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disorder;
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residues are 5-methylcytidines"
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                                                                                                   568; 132pp;
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== "2'-O-methoxyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      neurological disorder; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER
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Pred. No. 1.6e+03;
0; Mismatches 2
                                                                                                   English
                antisense oligonucleotide E2 synthase (mPGES-1). The e 9, more specifically to
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having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,

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RESULT 1489
ADM14481/c
ID ADM1448
AC ADM144B
AC ADM14AB
AC ADM14AB
AC ADM14AB
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADM14481 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:668
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                                                                                                                                                                                                                                                                                                                                                                   modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                              08-APR-2004.
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              (PHAA ) PHARMACIA CORP
                                                                                                                       25-SEP-2003; 2003WO-US030374
                                                                  25-SEP-2002; 2002US-0413549P
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/note= "phosphorothioate linkages and all cytidine
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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'note= "2',-O-methocyethyls"
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                                                                                                                                                                                                                                                                           _base= ОТНЬК
e= "2'-O-methoxyethyls"
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Pred. No. 1.
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RESULT 1490
ADM14501/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gierse JK;
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                modified_base
                                                                                                                            Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADM14501 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20
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                                                                                                                                                                                                       Synthetic
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                                                                                                                                                                                                                          sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                671 TGGCTCACTGCAACCTCTGC 690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mPGES-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rescreacrecascerecec 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chimeric antisense oligonucleotide SEQ ID NO:688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NO 668; 132pp; English.
         /*tag= b
/mod base= OTHER
/note= "phosphorothioate linkages
/note= "phosphorothioate linkages
residues are 5-methylcytidines"
                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; 5 C; 9
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Pred. No. 1
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Mismatches
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              chimeric; antisense oligonucleotide; phosphorothicate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory;
                                                                                                    Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1309.
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                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GTCTCGAACTCCTGGCCTCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 5 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunological, cardiovascular or neurological disorder
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                                                                                                                                                                                                                                                                           /*tag=
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                                                                                                                                                                                                                                                     /mod
                                                                                                                                                                                                                                                                                                                                       residues
                                                                                                                                                                                                                                                                                                                                                                      note= "phosphorothicate linkages"
                                                                                              'note=
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                                                                                                                                base= OTHER
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                                                                                              "2'-O-methoxyethyls"
                                                                                                                                                                                                                         "2'-0-methocyethyls'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                          are 5-methylcytidines"
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Pred.
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No. 1.
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Best Local
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                                                                                                                                                                                                     antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                      antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                  Sequence
                                                                                                                                                                                       ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PHAA ) PHARMACIA CORP
                         1001 CAAGCGATTCTCCTGTCTCA 1020
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                                                                          18; Conserv
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 CAAGCGATTCTCCCGCCTCA 1
                                                                                                                                                  20 BP; 4
                                                                                                                                                                                       immunological, cardiovascular or neurological disorder
                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ID NO 38; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2003WO-US030374
                                                                                                                                                  A; 3 C; 9
                                                                                           1.7%;
                                                                          0;
                                                                                                                                                  G; 4 T; 0 U; 0 Other;
                                                                                           Score 16.8;
Pred. No. 1.
                                                                            Mismatches
                                                                                             1.6e+03
                                                                                                               DB 1;
                                                                                                               Length 20
                                                                            Indels
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e.g.,

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RESULT 1495
ADM14695/c
ID ADM1469
 chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1 impES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
                                  Synthetic.
                                                                                        Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorcardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                               Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:882.
                                                                                                                                                                                                                                                                                                                  01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                      ADM14695;
                                                                                                                                                                                                                                                                                                                                                                                           ADM14695 standard; DNA;
                                                    Homo sapiens
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Location/Qualifiers
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WO2004028458-A2

NO:86

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ADM13899/
ID ADM1
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AC ADM1
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                                                                                                                                                                                                        RESULT 1496
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1 chimeric antisense oligonuclectides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; SEQ ID NO 882; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-SEP-2002; 2002US-0413549P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ophthalmic,
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                              01-JUL-2004
                                                                                           ADM13899;
                                                                                                                                                 ADM13899 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                    CCTGCCCCAGCCTCCTGAGT 731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 4 A; 4 C; 10 G; 2 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        immunological,
                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                              (first entry)
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16. .20
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residues are 5-methylcytidines"
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/note= "2'-O-methoxyethyls"
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/note= "2'-O-methocyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                      1.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                              Score 16.8; D
Pred. No. 1.6e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
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Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human mPGES-1 chimeric antisense oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                  modified_base
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                                                                                                                                               modified_base
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WO2004028458-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mmunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                  16. .20
                                                                                                                                                                                                                                                                                                 note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
                                                                                       /*tag= c
/mod_base= OTHER
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/note= "2'-0-methocyethyls"
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                                                           note=
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                                                           "2'-0-methoxyethyls'
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08-APR-2004

25-SEP-2003; 2003WO-US030374

25-SEP-2002; 2002US-0413549P (PHAA ) PHARMACIA CORP

Gierse JK;

WPI; 2004-305094/28

New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or e.g.,

Claim 4; SEQ ID NO 86; 132pp; English

The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mpCES-1). The CC thuman mpCES-1 gene is located on chromosome 9, more specifically to CC 9934.3. The present invention also describes: (1) antisense compounds, CC mpCES-1, which specifically hybridise with the nucleic acid encoding CC mpCES-1, which specifically hybridise with the nucleic acid encoding CC inhibits its expression; (2) a method of inhibiting the expression of CC draving a disease or condition associated with mpCES-1 and CC having a disease or condition associated with mpCES-1. MPGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antisinflammatory, neuroprotective, nouroprotective, antiarthritic, vasotropic, CC ophthalmological, immunomodulator, cardiant, neuroprotective and cardiovascular activities, and can be used as mpCES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpCES-1 e.g., inflammation, Alzheimer's CC ophthalmic, immunological, cardiovascular or neurological disorder.

Sequence 20 BP; 3 A. 6 C; 9 G; 2 T; 0 U; 0 Other;

¿Query Match

1.7%; Score 16.8; DB 1; Length 20,

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ADM14111/c
ID ADM14111
XX ADM1411
XX ADM1411
XX O1-JUL-
XX Chimeri
XW Chimeri
XW Chimeri
XW Inmuunon
XX Inmuunon
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromes persent synthase targeted on chromes of the present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encodi mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 an
                                                                                                                                                                                                                                                                            New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:298.
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                                                                                                                                                                                                  Claim 4; SEQ
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/note= "phosphorothioate linkages
residues are 5-methylcytidines"
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RESULT 1498
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Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inhibits its expression; (2) a method of inhibiting the expression mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chim antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunomodulator; cardiant; neuroprotective; antiinflammatory;
neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological;
immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chimeric; antisense oligonucleotide; phosphorothioate; human;
microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor;
microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:678
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                                                      25-SEP-2002; 2002US-0413549P
                                                                                  25-SEP-2003;
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                          (PHAA ) PHARMACIA CORP
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18; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                              disorder;
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                                                                                                                                                                                                                                                                                 /note= "phosphorothioate linkages residues are 5-methylcytidines"
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/note= "2'-O-methocyethyls"
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e= "2'-O-methoxyethyls"
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Pred. No. 1
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Gierse JK;

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RESULT 1499
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Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                        microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADM14603 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:790.
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                                                                                                                           residues
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                                                                                                                        /note= "phosphorothioate linkages residues are 5-methylcytidines"
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The present sequence represents a chimeric antisense oligonucleotide contargeted to human microsomal prostaglandin E2 synthase (mpCES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to 9344.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mpGES-1, which specifically hybridise, with the nucleic acid encoding CC mpGES-1 in cells or tissues; and (3) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, and can cophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition associated with mpGES-1 e.g., inflammation, Alzheimer's compound condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition in cardiovascular activities, and cardiovascular activities of condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition in cardiovascular activities of cardiovascula
RESULT 1500
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                chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
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microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mGGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or
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/note= "2'-O-methocyethyls"
16. .20
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/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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e= "2'-O-methoxyethyls"
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RESULT 1502
ADM15380/c
ID ADM1538
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ADM15380/c
ID ADM1538
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AC ADM1538
AC A
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1567
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                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                          25-SEP-2002; 2002US-0413549P
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18; Conserv
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larity 90.0%;
Conservative
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residues are 5-methylcytidines"
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'note= "2'-O-methocyethyls"
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e= "2'-O-methoxyethyls"
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Pred. No. 1.
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Claim 4;

SEQ ID NO 1567; 132pp; English

08-APR-2004.

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RESULT 1503
ADM14342/c
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Best Local &
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1. .20
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/note= "phosphorothioate linkages and
residues are 5-methylcytidines"
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/mote= "2'-O-methocyethyls"
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e= "2'-O-methoxyethyls"
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Pred. No. 1.
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L.6e+03;
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RESULT 1504
ADM14458/c
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Best Local Similarity
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                                                                                                                                                                       chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
       Key
modified_base
                                                                Homo sapiens.
Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                    ADM14458
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                        Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disor
                                                                                                                                                                                                                                                                                                          Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:645
                                                                                                                                                                                                                                                                                                                                                                                                                      ADM14458 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ophthalmic,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  resent sequence represents a chimeric antisense oligonucleotide ted to human microsomal prostaglandin E2 synthase (mPGES-1). The mPGES-1 pene is located on chromosome 9, more specifically to 3. The present invention also describes: (1) antisense compounds,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 5 A; 6 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunological, cardiovascular or neurological
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             Location/Qualifiers
1. .20
                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
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Pred. No. 1.6e+03;
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ADM13854/c
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                           having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1 mPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                                                                                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                         01-JUL-2004 (first entry)
                                                                                                      ADM13854 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                        ophthalmic,
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18; Conserv
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                                                                                                                                                                                                                                              Conservative
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residues are 5-methylcytidines"
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== "2'-O-methoxyethyls"
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                                                                                                                                                                                                                                                             Score 16.8;
Pred. No. 1.
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Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:41.

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The present sequence represents a chimeric antisense oligonucleotide crargeted to human microsomal prostaglandin E2 synthase (mpGES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1 in cells or tissues; and (3) a method of inhibiting the expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antistense oligonucleotides and antisense compounds have cytostatic, cc antidiabetic, immunomodulator, cardiant, neuroprotective, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
Query Match 1.7%;
Best Local Similarity 90.0%;
Matches 18; Conservative
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Score 16.8; DB 1;
Pred. No. 1.6e+03;
0; Mismatches
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                             The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9434.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of
                                                                                                                                                                                                                                                    Claim 4;
                                                                                                                                                                                                                                                                                                                             New antisense compound, having a sequence targeted to a nucleic avenceding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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  mPGES-1 in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                                                                                 SEQ ID NO 862; 132pp; English.
  cells or tissues; and (3)
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/note= "2'-
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/mod_base=
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residues are 5-methylcytidines"
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ADM14025/c
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antinflammatory, neuroprotective, notropic, antinflammatory, neuroprotective, notropic, antinflammatory, neuroprotective, notropic, antinflammatory, neuroprotective, notropic, antiserse compound be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpEES-1; mpEES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzhelmer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADM14025 standard; DNA;
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WPI; 2004-305094/28...
                              Gierse
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                                                                                                                                                                                                                                                                                                                  modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:212
                                                            (PHAA ) PHARMACIA CORP
                                                                                        25-SEP-2002; 2002US-0413549P
                                                                                                                    25-SEP-2003; 2003WO-US030374
                                                                                                                                                   08-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disorder; neurological disorder; ss.
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                  /note= "phosphorothioate linkages residues are 5-methylcytidines"
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mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                  mod_base= OTHER
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                                                                                                                                                                                                           "2'-0-methoxyethyls"
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Pred. No. 1.
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antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                           having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1 MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compound, having a sequence targeted to a nucleic encoding mPGES-1, useful for preparing a composition for treatinflammation, Alzheimer's disease, arthritis, diabetes, cancer
Sequence 20 BP; 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4;
                                              ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                            human
                                                                                                                                                                                                                                                                                                                                                                                                                                         mPGES-1
                                                                                                                                                                                                                                                                                                                                                                                                                 ent sequence represents a chimeric antisense oligonucleotide to human microsomal prostaglandin E2 synthase (mPGES-1). The GBS-1 gene is located on chromosome 9, more specifically to The present invention also describes: (1) antisense compounds,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 212; 132pp; English
                                           immunological, cardiovascular or neurological disorder
A; 8 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a nucleic
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RESULT 1508
ADM14469/c
ID ADM1446
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Best Local Similarity
Matches 18; Conserv
  ADM14469 standard; DNA; 20 BP
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                                                                      AGTGCTGGGATTACAGGCGT 410
                                                       AGTGCTGGGATGACAGGCAT 1
                                                                                              1.7%;
ilarity 90.0%;
Conservative
                                                                                              ٥,
                                                                                                         Score 16.8;
Pred. No. 1.
                                                                                               Mismatches
                                                                                                          6e+03
                                                                                                                    BB
                                                                                                                   1;
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                                                                                                Indels
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ADM14469;

(first entry)

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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpCES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; secardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                                                                                                                                                                      Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:656
              modified_base
                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                   Homo
                                                                           modified_base
                                                                                                                                                    modified_base
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/mod_base= OTHER
/note= "2'.-O-methocyethyls"
16. 20
/*tag= c
                                                                              residues are
                                                                                                                                                                    Location/Qualifiers
                                                                                           /note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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RESULT 1509
ADM14642/c
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Best Local S
Matches 18
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                                                                chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         targeted to human mPGES-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (PHAA ) PHARMACIA CORP
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Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorcardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                  Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:829.
                                                                                                                                                                                                                                                                                                                                                                                  ADM14642 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ophthalmic,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GGCTCACTGCAGCCTCCGCC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence represents a chimeric antisense oligonucleotide human microsomal prostaglandin E2 synthase (mPGES-1). The control on chromosome 9, more specifically to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunological,
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                                                                                                                                                                                                                                                                                 (first entry)
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/note= "2'.-O-methoxyethyls"
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Query Match
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pred. No. 1.6e+03
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RESULT 1510 ADM14763/c ID ADM14763

standard; DNA; 20

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective.
antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                     New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor;
prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
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ADM14262/c
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chimeric; antisense oligonucleotide; phosphorothioate; human;
microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor;
microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                      New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:449.
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                     Claim 4; SEQ
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                                                                                                      WPI; 2004-305094/28.
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                    ID NO 449; 132pp; English
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/note= "phosphorothicate linkages and all cytidine
residues are 5-methylcytidines"
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e= "2'-O-methoxyethyls"
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Pred. No. 1.
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The present sequence represents a chimeric antisense oligonucleotide

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RESULT 1512
ADM14410/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonuclectides and antisense compounds have cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The
                                                                                                                                                                                                                                                                                                                        modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorcardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:597
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
              25-SEP-2003; 2003WO-US030374
                                                                                                                                                                     modified_base
                                                                                                                                                                                                                                                                                                                                                                                                    Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological;
                                                    08-APR-2004.
                                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunomodulator; cardiant; neuroprotective; antiinflammatory;
                                                                                     WO2004028458-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                    sapiens.
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18; Conser
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                                                                                                                                                                                                                                                              residues are
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                                                                                                                                                                                                                                                            'notē= "phosphorothioate linkages and all cytidine
:esidues are 5-methylcytidines"
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e= "2'-O-methocyethyls"
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                                                                                                                                                                                                                                                                                             base= OTHER
                                                                                                                      base= OTHER
== "2'-O-methoxyethyls"
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Pred. No. 1.6e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 U;
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RESULT 1513
ADM14596/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents a chimeric antisense oligonucleotide crargeted to human microsomal prostaglandin E2 synthase (mpGES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1 in cells or tissues; and (3) a method of thibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or CC conthibulatic, immunofical acradiance reperfusion injury, or CC conthibulatic, diabetes, cancer, ischaemia or reperfusion injury, or CC conthibulatic, diabetes, cancer, ischaemia or reperfusion injury, or contribution immunofical acradiance compound cardiovascular activities.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 18
                                                                                                                                                           Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disordecardiovascular disorder; neurological disorder
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                                                                                                                                                                                                                       immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                  chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; SEQ ID NO 597; 132pp; English.
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                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                   Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:783.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20
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                                                                                                                   Synthetic
                                                                                                                                   Homo sapiens.
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                                                                          Location/Qualifiers
                                  /*tag=
note= "phosphorothicate linkages"
                      mod_base= OTHER
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Pred. No. 1.
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      and all cytidine
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RESULT 1514
ADM14660/c
ID ADM1466
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AC ADM1466
XC ADM1466
XC ADM1466
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DT 01-JUL-
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DE Human n
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KW chimeri
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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 chimeric; antisense oligonucleotide; phosphorothioate; human;
microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor;
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a chimeric antisense oligonuclectide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9034.3. The present invention also describes: (1) antisense compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; SEQ ID NO 783; 132pp; English.
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                                                    Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:847.
                                                                                         01-JUL-2004
                                                                                                                                                              ADM14660 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                           ophthalmic,
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                                                                                                                                                                                                                                                                                                                                                                                                                         immunological, cardiovascular or neurological disorder.
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                                                                                         (first entry)
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16. .20
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/note= "2'-O-methocyethyls"
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                                                                                                                                                                                                                                                                                                                                   Score 16.8; DB 1;
Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                       1;
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1056 CCACACCCCGCTAATTTTTG 1075

a. 1,

Query Match Best Local Matches

18;

Conservative

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Similarity

1.7%;

Score 16.8; Pred. No. 1 Mismatches

1.6e+03 DB 1;

Length 20 Indels

<u>.</u>

Gaps

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to C2 9344.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding confidence of thibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal chaving a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic, cantisense oligonucleotides and antisense compounds have cytostatic, ophthalmological, immunomodulatory, and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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neuroprotective; nootropic; antlarthritic; vasotropic; ophthalmological;
immunomodulatory; cardiovascular; gene therapy; inflammation;
Alzheimer's disease; arthritis; diabetes; cancer; ischaemia;
reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gierse JK;
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Synthetic.
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Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                              ischemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (PHAA ) PHARMACIA CORP
 BP; 7 A; 2 C; 6 G; 5 T; 0 U; 0 Other;
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residues are 5-methylcytidines"
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ADM14676/c
                  The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9344.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibite its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; contactic; antidiabetic; immunomodulator; cardiant; neuroprotective; antinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
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                                                                                                                                                                                                                                                                                                                                                      New antisense compound, having a sequence targeted to a encoding mPGES-1, useful for preparing a composition for inflammation, Alzheimer's disease, arthritis, diabetes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:863
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                                                                                                                                                                                                                                                                              Claim 4;
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esidues are 5-methylcytidines"
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  compounds have cytostatic,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischammia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammaror.
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New antisense compound,

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RESULT 1517
ADM14269/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 934.3. The present invention also describes: (1) antisense compounds, process, which specifically hybridise with the nucleic acid encoding medss-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of medss-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with medss-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cycostatic, antisense oligonucleotides and antisense compounds have cycostatic, antisense oligonucleotides and antisense compounds have cycostatic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmological, disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmological contents of the condition and contents of the condition associated with mPGES-1 e.g., inflammation, Alzheimer's contents of the condition associated with mPGES-1 e.g., inflammation or reperfusion injury, or ophthalmological contents of the condition associated with mPGES-1 e.g., inflammation, Alzheimer's contents of the condition and condition associated with mPGES-1 e.g., inflammation, Alzheimer's contents of the condition and condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition associated with mPGES-1 e.g., inflammation, Alzheimer's conditions and conditions anamed conditions and conditions and conditions and conditions and 
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antifilammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADM14269
                                                                                                modified_base
                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                              Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorcardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:456.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunological, cardiovascular or neurological disorder.
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/*tag= c
/mod_base= OTHER
/mote= "2'-O-methoxyethyls"
                                                                                                  /note= "
16. .20
                                                                                                                                                                                                                      residues
                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                    /note= "phosphorothioate linkages and residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                       *tag=
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e= "2'-O-methocyethyls"
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.6e+03
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                                                                                                                                                                                                                                          all cytidine
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The present sequence represents a chimeric antisense oligonucleotide cargeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to compare the present invention also describes: (1) antisense compounds, of the present invention also describes: (1) antisense compounds, or mpGES-1, which specifically hybridise with the nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding mpGES-1 in cells or tissues, and (3) a method of inhibiting the expression of mpGES-1 in cells or tissues, and (3) a method of treating an animal caning a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonuclectides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiarthritic, vasotropic, antininflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, cophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                               Query Match
                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; SEQ ID NO 456; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                          Sequence 20
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                                                              Local
1004 GCGATTCTCCTGTCTCAGCC 1023
                                           18;
                                                              Similarity
                                                                                                                          BP; 4 A; 4 C; 10
                                           Conservative
                                                              1.7%;
                                           <u>.</u>
                                                              Score 16.8;
Pred. No. 1
                                                                                                                          G; 2 T; 0 U;
                                             Mismatches
                                                                1.6e+03
                                                                                                                             0 Other;
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                                           2
                                                                                 Length
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RESULT 1518 ADM14328/c ID ADM1432 S 밁 chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; noctropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; ADM14328 standard; DNA; 20 Human 01-JUL-2004 ADM14328; 20 (first entry) chimeric antisense oligonucleotide SEQ ID ВP NO:515 <u>,</u> Gaps

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RESULT 1519
ADM14470/c
ID ADM1447
XX
AC ADM1447
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                                                                                                                                                                 Query Match
Best Local S
Matches 18
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encoding mPGES-1, useful for preparing a composition for treating
inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4;
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    ADM14470
                               ADM14470 standard;
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                                                                                                                                     708
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                                                                                                                                                                   18;
                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ
                                                                                                                                                                                                                            20
                                                                                                                              TTCTCCTGCCCCAGCCTCCT 727
                                                                                                          TTCTCCCGCCTCAGCCTCCT 1
                                                                                                                                                                                                                            BP; 6 A;
                                                                                                                                                                   Conservative
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16. .20
/*tag=
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residues are 5-methylcytidines"
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                                 DNA;
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e= "2'-O-methocyethyls"
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== "2'-O-methoxyethyls"
                                                                                                                                                                               90.0%;
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                                 ВP
                                                                                                                                                                   0
                                                                                                                                                                               Score 16.8;
Pred. No. 1.
                                                                                                                                                                                                                             G; 1 T; 0
                                                                                                                                                                   Mismatches
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                                                                                                                                                                                 6e+03
                                                                                                                                                                                              DB 1; Length 20;
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                                                                                                                                                                                                                               Other;
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                                                                                                                                                                    Gaps
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microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological;
                                                                                                                                                                                                                                                                                                                   25-SEP-2003; 2003WO-US030374
                                                                                                                                                                                                                                                                                                                                      08-APR-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      chimeric; antisense oligonucleotide; phosphorothioate;
microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kuman mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:657
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mmunomodulatory; cardiovascular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disorder;
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/note= "2'-
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                              note= "phosphorothioate linkages
residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                            OTHER
-O-methocyethyls"
                                                                                                                                                                                                                                                                                                                                                                                   OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         and
                                                                                                                                                                                                                                                                                                                                                                                                                                                          all cytidine
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New antisense compound, having a sequence targeted to a nucleic acid encoding mPGBS-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or e.g.,

Gierse

WPI; 2004-305094/28

25-SEP-2002; 2002US-0413549P

(PHAA ) PHARMACIA CORP

Claim 4; SEQ ID NO 657; 132pp; English

The present sequence represents a chimeric antisense oligonucleotide cargeted to human microsomal prostaglandin E2 synthase (mpGES-1). The C1 human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, C2 having a sequence comprising 8-30 bp targeted to a nucleic acid encoding captured by the specifically hybridise with the nucleic acid encoding captured in thibits its expression; (2) a method of inhibiting the expression of captured captured conditions associated with mpGES-1 and captured captur immunological, cardiovascular or neurological

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 RESULT 1520
ADM15246/c
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20
                                                                                      New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.cinflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADM15246;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADM15246 standard; DNA;
          The present targeted to
                                                   Claim 4;
                                                                                                                                           WPI; 2004-305094/28.
                                                                                                                                                                                                                       25-SEP-2002; 2002US-0413549P
                                                                                                                                                                                                                                                 25-SEP-2003; 2003WO-US030374
                                                                                                                                                                                                                                                                                                   WO2004028458-A2
                                                                                                                                                                                                                                                                                                                                                              modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                              (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1057 CACACCCCGCTAATTTTTGT 1076
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                   SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CATACCCAGCTAATTTTTGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 8
sequence represents a chimeric antisense oligonucleotide human microsomal prostaglandin E2 synthase (mPGES-1). The -1 gene is located on chromosome 9, more specifically to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                 ID NO 1433; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                  16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "phosphorothioate linkages
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                         /*tag= c
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                              note=
                                                                                                                                                                                                                                                                                                                                                                             note=
                                                                                                                                                                                                                                                                                                                                                                                            mod base=
                                                                                                                                                                                                                                                                                                                                                                                                       *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                         mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Α,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 C; 5
                                                                                                                                                                                                                                                                                                                             "2'-0-methoxyethyls'
                                                                                                                                                                                                                                                                                                                                                                             "2'-0-methocyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ₽₽
                                                                                                                                                                                                                                                                                                                                                                                            OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 16.8; DB 1;
Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                             and all cytidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                   e.g.,
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RESULT 1521
ADM15325/c
ID ADM1532
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC having a sequence comporising 8-30 bp targeted to a nucleic acid encoding cc mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and cinhibits its expression; (2) a method of inhibiting the expression of cc mpGES-1 in cells or tissues; and (3) a method of treating an animal cc having a disease or condition associated with mpGES-1. MpGES-1 chimeric cantisense oligonucleotides and antisense compounds have cytostatic, cc antification immunomodulator, cardiant, neuroprotective, vasotropic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, cophthalmological, immunomodulatory and cardiovascular activities, and can compuse the used as mpGES-1 inhibitors and in gene therapy. The antisense compound cc an be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's composition associated with mpGES-1 e.g., inflammation, Alzheimer's cophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                     chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 5 A; 5 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADM15325 standard;
                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
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                                                                                                                WO2004028458-A2
                                                                                                                                                                                                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   213 GGTCTCGAACTCCCGACCTC 232
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GGTCTCGAACTCCTGGCCTC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                       /*tag= c
/mod_base= OTHER
/note= "2'-O-methoxyethyls"
                                                                                                                                                                                        /*tag= a
/mod_base= OTHER
/note= "2'-O-methocyethyls"
                                                                                                                                                                                                                                                                                /note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                   mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                        *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90.0%;
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
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25-SEP-2002; 2002US-0413549P 25-SEP-2003; 2003WO-US030374

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RESULT 1522
ADM15564/c
ID ADM1556
XX
AC ADM1556
XX
DE Human m
XX
DE Human m
XX
Chimeri
KW microsec
KW microsec
KW microsec
KW immunor
KW immunor
KW reperfi
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antiinflammatory, neuroprotective, nootropic, antiinflammatory, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibit its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             encoding mPGES-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                      chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                                                                                                                                                                                                    immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-JUL-2004
                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                    cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADM15564;
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         modified_base
                                                                                                                                                     modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2004-305094/28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGTTCACCATGTTCGCCAGG 812
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGTTCACCATGTTGCCCAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 5 A; 6 C; 6 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chimeric antisense oligonucleotide SEQ ID NO:1751.
                                                                                                        1. ...
/*tag= b
                                                                                                                                                                                                                                                                                                                                          disorder;
      residues
               /mod_base= ОТНЬК
/note= "phosphorothioate linkages
residues are 5-methylcytidines"
                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           132pp;
                                                                                                                                                                                                                                                                                                                                       neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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Pred. No. 1.
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                                                                     and
                                                                     all cytidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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Human mPGES-1 01-JUL-2004

chimeric antisense oligonucleotide SEQ ID

NO:109

(first entry)

chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory;

RESULT 1523 ADM13922/c

ADM13922 standard;

DNA;

0

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CC human mpGES-1 gene is located on chromosome 9, more specifically to CC 934.3. The present invention also describes: (1) antisense compounds, CC 934.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC mpGES-1 in cells or condition associated with mpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antinflammatory, neuroprotective, noctropic, antistflammatory, neuroprotective, noctropic, antistflammatory, immunomodulator, and in gene therapy. The antisense compound CC can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's compound contraction of the condition injury, or condition associated with mpGES-1 e.g., inflammation, Alzheimer's compound contraction injury, or contraction in contraction injury, or contraction in contraction in contraction injury, or contraction in contraction injury, or contraction in contraction in contraction injury, or contraction in contraction
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                                             S
                                                                                             Matches
                                                                                                                    Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic avenceding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gierse JK;
                                                                                                                                                                                                                                              disease, ar ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; SEQ ID
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                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PHAA ) PHARMACIA CORP.
                                                369
20
                                                                                                18;
                                                                                                                       Similarity
                                                  TCCACCTGCCTCAGCCTCCC 388
                                                                                                                                                                                              20
                                                                                                                                                                                              B₽;
                                                                                                                                                                                                                                           immunological, cardiovascular or neurological
                                                                                                Conservative
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e= "2'-O-methocyethyls"
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                                                                                                                         Score 16.8; DB 1;
Pred. No. 1.6e+03;
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                                                                                                                                                                                                   0 Other;
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CC targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The CC human mPGES-1 gene is located on chromosome 9, more specifically to CC 9934.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mPGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mPGES-1. MPGES-1 chimeric CC antisense oligonuclectides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, and can complaint activities, and can CC be used as mPGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or contition associated with mPGES-1 e.g., inflammation, Alzheimer's CC conthibalmic immunological, cancer, ischaemia or reperfusion injury, or conthibalmic immunological acradiant, schaemia or reperfusion injury, or conthibalmic immunological cardiant is schaemia or reperfusion injury, or conthibalmic immunological acradiant is schaemia or reperfusion injury, or conthibalmic immunological cardiant is schaemia or reperfusion injury, or conthibalmic immunological cardiant acradiant or continuological disorder.
                                                                                                                      Best Local Similarity Matches 18; Conserv
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                                                                                                                                                                                                                                                    Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                    1002 AAGCGATTCTCCTGTCTCAG 1021
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                                                                                                                                                                                                                                                BP; 4 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
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                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                               immunological,
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No. 1.
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                                                                                  The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                               antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 333; 132pp; English.
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prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
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residues are 5-methylcytidines"
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neuroprotective,
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   nootropic,
      antiarthritic,
   vasotropic,
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RESULT 1525
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Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:861.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUL-2004
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  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                   25-SEP-2002; 2002US-0413549P
                                                                                                                                                                                                                                                                                                                                       25-SEP-2003; 2003WO-US030374
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                          Gierse
                                                                                                                   WPI; 2004-305094/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 682 AACCTCTGCCTCCCGGGTTC 701
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                                                                                                                                                                                                                              PHARMACIA CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "phosphorothioate linkages and
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                _base= OTHER
.e= "2'-O-methoxyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "2'-0-methocyethyls"
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Pred. No. 1.6e+03;
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); Mismatches
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RESULT 1526
ADM14776/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                                                                                                                                                                                                                                                                                                        chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Çlaim 4; SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADM14776 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:963
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20
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                                                                                modified_base
                                                                                                                                                                                                                                     modified_base
                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                          Homo sapiens.
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                                                                                                                                                   modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                  /*tag=
                                                                                                                                                                       residues
                                                                                                                    /*tag= a
/mod_base=
                                                                                                                                                                       /note= "phosphorothioate linkages and
residues are 5-methylcytidines"
                                  'mod_base= OTHER
'note= "2'-O-methoxyethyls'
                                                                                                    note=
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                                                                                      . 20
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                                                                                                                                                                                                         base= OTHER
                                                                                                    "2'-0-methocyethyls"
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                                                     OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No. 1.60); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20;
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                                                                                                                                                                                         all cytidine
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WO2004028458-A2

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RESULT 1527
ADM14800/c
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGBS-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
             Homo sapiens.
Synthetic.
                                                                 Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorcardiovascular disorder; neurological disorder; ss.
                                                                                                                                        microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antifilammatory; neuroprotective; antioxidator; cardiant; neuroprotective; antioxidator; cardiant; neuroprotective; ophthalmological;
                                                                                                                                                                                                                     chimeric; antisense oligonucleotide; phosphorothioate; human;
                                                                                                                                                                                                                                                          Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:987.
                                                                                                                                                                                                                                                                                              01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                       ADM14800 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-305094/28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-SEP-2002; 2002US-0413549P.
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                                                                                                                                                                                                                                                                                                                                   ADM14800;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PHAA ) PHARMACIA CORP
                                                                                                                           mmunomodulatory; cardiovascular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           resent sequence represents a chimeric antisense oligonucleotide ted to human microsomal prostaglandin E2 synthase (mPGES-1). The mPGES-1 gene is located on chromosome 9, those specifically to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 963; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GCAGCCTCCGCCTCCCGGGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GCAACCTCTGCCTCCCGGGT 699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 3 A; 6 C; 10 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immunological, cardiovascular or neurological
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                                                                                                                                                                                                                                                                                            (first
                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90.0%;
                                                                                                                                                                                                                                                                                                                                                                       20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 16.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                         gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .6e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20
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                                                                                      disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disorder
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RESULT 1528
ADM14814/c
ID ADM1481
XX
AC ADM1481
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AC ADM1481
XX
OT 01-JUL-

ADM14814;

ADM14814 standard; DNA; 20 BP

01-JUL-2004

(first entry)

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Best Loc
Matches
                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense compound, having a sequence targeted to a nucleic avenceding mPGES-1, useful for preparing a composition for treating encoding mPGES-1, disease, arthritis, diabetes, cancer or inflammation, Alzheimer's disease, arthritis, diabetes, cancer or ischamia
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CACCATACCCAGCTAATTTT
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microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chimeric; antisense oligonucleotide; phosphorothioate;
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The present sequence represents a chimeric antisense oligonucleotide cc targeted to human microsomal prostaglandin E2 synthase (mpDES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antistabetic, immunomodulator, cardiant, neuroprotective, cc antistiflammatory, neuroprotective, noctropic, antistiflammatory, neuroprotective, noctropic, antistiflammatory, neuroprotective, noctropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's ophthalmic, immunological, cardiovascular or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.

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Matches 18
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           The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds
                                                                                                          New antisense compound, having a sequence targeted to a encoding mPGES-1, useful for preparing a composition for inflammation, Alzheimer's disease, arthritis, diabetes,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
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/mod_base= OTHER
/note= "2'-O-methoxyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
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residues are 5-methylcytidines"
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Pred. No. 1.6e+03;
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The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to CC dy34.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mPGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mPGES-1. MPGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, cantisnse oligonucleotides and antisense compounds have cytostatic, continifiammatory, neuroprotective, nootropic, antiarthritic, vasotropic, antiinflammatory, neuroprotective, nootropic, antiarthritics, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's CC ophthalmic, immunological, cardiovascular or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compound, having a sequence targeted to a nucleic avenceding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                       Sequence 20 BP; 2 A; 4 C; 12 G; 2 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; SEQ ID NO 1713; 132pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
     DB 1; Length 20;
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RESULT 1531
ADO46482
        TO A CANA WAX WAX SO XX CAN A 
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                                                                                   23-APR-2002;
23-APR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; 88; interleukin-4 receptor; IL-4; interleukin-5 receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human oligonucleotide #1848.
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                                                                                                                                                                                                                                                                                                                                                                                     US2004049022-A1
(NYCE/) NYCE J W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       370 CCACCTGCCTCAGCCTCCCA 389
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Similarity 90.0%;
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                                                                                   2002WO-US013135.
2002WO-US013143.
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region codon, coding region, 5' or 3' intron-exon junction, intron or region codon, coding region, 5' or 3' intron-exon junction, intron or region codon, coding region of a nucleic acid target cohosen from a gene encoding interleukin (IL) 4 receptor, interleukin (IL) 6.5 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDB4 A, PDB4 B, PDB4 C or PDB4 D. The invention code or management of screening a candidate compound that binds to come or more nucleic acid target(s) or expressed product(s), for the code or or more nucleic acid target(s) or expressed product(s), for the code or or management of a respiratory or lung disease. The code or management of a receptor, interleukin-5 receptor, CC craj, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDB4 B, PDB4 B, PDB4 C, or PDB4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The complete of increased levels of, adenosine and/or levels of adenosine A CC receptor(s), and/or asthma and/or lung allergies associated with cc inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, constitute pulmonary disease (COPD), calleric thiritis article prespiratory of disease (COPD), calleric thiritis article prespiratory of disease sendones. Disease (COPD)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
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(TANG/) TANG L.
(AGUI/) AGUILAR D.
(MILL/) MILLER S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
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lung disease; hyper-responsiveness; adenosine; adenosine; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic chronic obstructive pulmonary disease; COPD; allergic rhi
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                                                              Human; 88; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor;
                                                                                                                                                                  Human oligonucleotide
                                                                                                                                                                                                        15-JUL-2004
                                                                                                                                                                                                                                                                                     ADO44692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bronchoconstriction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction o
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                                                                                                                                                                                                                                                                                                                                                                                                                             535
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MILLER S.
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LU H.
                                                                                                                                                                                                                                                                                     standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                     CTCCCACCTCAGCCTCCCAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                             CTCCTGCCTCAGCCTCCCAA 554
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in S, Lu H,
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                                                                                                                                                                                                        (first entry)
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Cong
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90.0%;
                                                                                                                                                                  #58.
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ng H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                           cystic fibrosis; CF;
          rhinitis;
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The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region CC with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target CC chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-6. receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to CC one or more nucleic acid target(s) or expressed product(s), for the CC prevention and/or treatment of a respiratory or lung disease. The CC oligonucleotides are useful for reducing or inhibiting expression of a CC gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CC CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC useful for preventing or treating a respiratory or lung disease. The CC respiratory or lung disease is associated with hyper-responsiveness to inflammation or an inflammatory disease. The receptor(s), and/or asthma and/or lung allergies associated with CC is chosen from airway inflammation, allergy, asthma, impeded respiration, CC cystic fibrosis (CF), chronic obstructive pulmonary disease (COPD), alleric respirators on inflammation, allergy, asthma, impeded respiration, CC cystic fibrosis (CF), chronic obstructive pulmonary disease (COPD)
                                                                                                        Matches
                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nyce JW, Sa
Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (NYCE/)
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23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 2; SEQ ID NO 58; 174pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            asthma.
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                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                       allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the
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MILLER S.
SHAHABUDDIN
LU H.
CONG H.
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RESULT 1533 ADO46473

ADO46473 standard; DNA;

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The invention relates to oligonucleotides anti-sense to an initiation CC codon, coding region, 5' or 3' intron-exon junction, intron or region CC with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target CC chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL)-6. Freceptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, DE4A A, PDE4 B, PDE4 C or PDE4 D, The invention CC also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the CC prevention and/or treatment of a respiratory or lung disease. The CC prevention and/or treatment of a receptor, interleukin-5 receptor, CC criptase b, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for reducing or inhibiting expression of a C gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CC criptase b, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The C useful for preventing or treating a respiratory or lung disease. The C creptor(s), and/or asthma and/or lavels of adenosine A C receptor(s), and/or asthma and/or lavels associated with C inflammation or an inflammatory disease. The respiratory or lung disease (C is chosen from airway inflammation, allergy, asthma, impeded respiration, C cystic fibrosis (CF), chronic obstructive pulmonary disease (COPD), chronic obstructive pulmonary disease.
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               allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 2;
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Shahabuddin
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23-APR-2002; 2002WO-US013143.
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in S, Lu H,
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RESULT 1534
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                        The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (IJ)-4 receptor, interleukin (II-5 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention also relates to a method of screening a candidate compound that binds to
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23-APR-2002;
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  elates to a method of screening a candidate compound that bin more nucleic acid target(s) or expressed product(s), for the
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Similarity 90.0%;
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CC prevention and/or treatment of a respiratory or lung disease. The CC oligonucleotides are useful for reducing or inhibiting expression of a CC gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CC CCRI, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC useful for preventing or treating a respiratory or lung disease. The CC respiratory or lung disease is associated with hyper-responsiveness to CC and/or increased levels of, adenosine and/or levels of adenosine A CC receptor(s), and/or asthma and/or lung allergies associated with CC inflammation or an inflammatory disease. The respiratory or lung disease in flammatory disease is associated with a ccompany or lung disease confictive particles associated with a ccompany or lung disease confictive particles associated with a ccompany or lung disease confictive particles as syndrome, pulmonary construction or an inflammation, allergy, asthma, impeded respiration, constitutive particles as syndrome, pulmonary construction or allergic rhinitis, acute respiratory distress syndrome, pulmonary construction, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the invention.

Sequence 20 BP; 3 A; 7 C; 4 G; 6 T; 0 U; 0 Other;

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RESULT 1535
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                 Human; 88; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; asthma; lung allergy; inflammation; inflammatory disease; circular allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension;
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                              Nyce JW, Sandrasagra A,
Shahabuddin S, Lu H, C
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23-APR-2002;
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                                                                                                                                       (NYCE/) NYCE J W. (SAND/) SANDRASAG
                                                                                           SHAH/)
                                                                                                                                                                                                                                                                                                        inflammation; bronchitis; airway obstruction; bronchoconstriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           932 TCACTCTGTTACCCAGGCTG 951
                                                                                                                                                                                                                                                                                                                                                                                                                            oligonucleotide #1804.
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                                                                     CONG 1
                                                                                                                             TANG
                                                                                            SHAHABUDDIN
                                                                                                      MILLER S.
                                                                                                                 AGUILAR D.
                                                                                                                                        SANDRASAGRA A.
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                                                                                                                                                                          2002WO-US013135.
2002WO-US013143.
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ong H;
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Pred. No. 1.6e+03;
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                                             Aguilar
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Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, PANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g. asthma.

Claim 2; SEQ ID ŏ 1805; 174pp; English.

chosen from a gene encoding interleukin (II)-4 receptor, CRM, VCAM, VCAM codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region bronchoconstriction. This sequence represents an oligonucleotide

Sequence 20 BP; 4 A; σ C; 7 G; 3 T; 0 U; 0 Other;

Ś Query Match Best Local S Matches 18 1033 GCTGGGATTACGGGCACCTG 1052 18; Similarity Conservative 1.7%; 0 Score 16.8; Pred. No. 1 Mismatches .6e+03 DB 1; Length 20; Indels 0 Gaps 0

RESULT 1536 뭉 GCTGGGATTACAGGCACCCG 20

Human oligonucleotide #1798 ADO46432 standard; DNA; (first 20

Human, ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Botaxin-1; RANTES; MCP4; CD33; ICAM; tryptase a; tryptase b; PD84 A; PD84 B; PD84 C; PD84 D; respiratory disease; lung disease; hyper-responsiveness; adenosine, adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; alivay inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction. 23-APR-2002; 23-APR-2002; 11-MAR-2004 US2004049022-A1 25-JUL-2003; 2002WO-US013135 2002WO-US013143 2003US-00627930

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                 Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         is chosen from airway inflammation, allergy, asthma, impeded responses to the fibrosis (CF), chronic obstructive pulmonary disease (COPD allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction of bronchoconstriction. This sequence represents an oligonucleotide of the control o
                                                                                                                                                                                                                                                                                                                                       ADO46461 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel single or multiple target oligonucleotide anti-sense to e.g. Conitiation codon, intron of respiratory disease-relevant gene e.g. Con RANTES, MCP4, useful for prophylaxis or treating respiratory disease
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MILLER S.
SHAHABUDDIN S.
LU H.
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SANDRASAGRA
TANG L.
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ong H;
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airway inflammation; allergy; impeded respiration; cystic fichronic obstructive pulmonary disease; (CDP); allergic rhinit acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoco
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                                                                                                                                                                                                                                                         Claim
                                                                                                                                                                                                                                                                                   Movel single or multiple target oligonucleotide anti-sense to initiation codon, intron of respiratory disease-relevant gene RANTES, MCP4, useful for prophylaxis or treating respiratory or
                                                                                                                                                                                                                                                                                                                                           Shahabuddin S,
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                                                                                                                                                                                                                                                                                     respiratory disease
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gene e.g. CCR1,
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CC chosen from a gene encoding interleukin (II).4 receptor, interleukin (II) CC -5 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to CC one or more nucleic acid target(s) or expressed product(s), for the CC prevention and/or treatment of a respiratory or lung disease. The CC prevention and/or treatment for reducing or inhibiting expression of a CC gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CC CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC useful for preventing or treating a respiratory or lung disease. The CC respiratory or lung disease is associated with hyper-responsiveness to CC and/or increased levels of, adenosine and/or levels of adenosine A CC receptor(s), and/or asthma and/or lung allergies associated with CC inflammation or an inflammation, allergy, asthma, impeded respiration, CC cystic fibrosis (CP), chronic obstructive pulmonary disease (COPD), CC allergic rhintline acute respiratory distease surdarea envalvance of allergic rhintline.

allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction obronchoconstriction. This sequence represents an oligonucleotide

Sequence

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0 Other;

Query Match Best Local S Matches 18 936 l Similarity 18; Conserv TCTGTTACCCAGGCTGGAGT 955 larity 90.0%; Conservative 0 Score 16.8; Pred. No. 1 Mismatches 1.6e+03 B 1; <u>ب</u> Length 0 Gaps 0

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RESULT 1538
ADO46483
                                           CC codon, coding region, 5 or 3 intron-exon junction, intron or region CC with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target C chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL) CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to CC one or more nucleic acid target(s) or expressed product(s), for the CC prevention and/or treatment of a respiratory or lung disease. The CC oligonucleotides are useful for reducing or inhibiting expression of a CC CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a CC crespiratory or lung disease. The CC crespiratory or lung disease. The CC crespiratory or lung disease is associated with hyper-responsiveness to CC and/or increased levels of, adenosine and/or levels of adenosine A CC crecptor(s), and/or asthma and/or lung allergies associated with CC inflammation or an inflammatory disease. The respiratory or lung disease
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initiation codon, intron of respiratory disease-relevant gene e-
RANTES, MCP4, useful for prophylaxis or treating respiratory dis
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Shahabuddin
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23-APR-2002;
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is chosen from airway inflammation, allergy, asthma, impedecystic fibrosis (CF), chronic obstructive pulmonary disease allergic rhinitis, acute respiratory distress avadrama and
                                                                                                                                                                                                                                                                                                                                                                                   Claim 2;
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CCR1; CCR3; Ectaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a;
tryptase b; DD4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease;
lung disease; hyper-responsiveness; adenosine; adenosine A recepto
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TANG L.
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ong H;
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                                                                                                                                                                                                                                                                                                                                                                                   English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Miller
                     impeded respiration
disease (COPD),
                                                                                                                                                                                                                                                                                                                                                                                                                                       disease
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RESULT 1539
AD04526
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Ebtaxin-1; RANTES; MCP4; CD23; ICAM; CCRM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; CDP; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADO45260 standard;
                                                                                                                                                                                                                                                Novel single or multiple target oligonucleotide anti-sense to e.g. coinitiation codon, intron of respiratory disease-relevant gene e.g. Coronitiation codon, intron of respiratory disease relevant gene e.g. Coronitiation (CP4, useful for prophylaxis or treating respiratory disease
                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-293804/27.
                                                                                                                                                                                                                                                                                                                                                                                                     Nyce JW, Sandrasagra
Shahabuddin S, Lu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-APR-2002; 2002WO-US013135.
23-APR-2002; 2002WO-US013143.
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MILLER S.
SHAHABUDDIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CONG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TANG L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SANDRASAGRA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
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mg H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  'n
                                                                                                                                                                                                                                                                                                                                                                                                                                       Miller
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The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3' end of a nucleic acid target chosen from a gene encoding interleukin (IL)-4 receptor, interleukin -5 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention

target

(II)

CCR1, se e.g

Claim 2; SEQ ID NO 626; 174pp; English

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RESULT 1540
ADO46435
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Best Local S
Matches 18
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        Shahabuddin
                                             Nусе JW,
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23-APR-2002; 2002WO-US013143
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                                                                                                                    SHAHABUDDIN
LU H.
CONG H.
                                                                                                                                                                                                                                        SANDRASAGRA
TANG L.
AGUILAR D.
MILLER S.
                                                                                                                                                                                                                                                                                                                                                                                                              NYCE J W.
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Sandrasagra A, Tang L,
in S, Lu H, Cong H;
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Pred. No. 1.
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                                         Aguilar D,
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                                             Miller
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Novel single or multiple target oligonucleotide anti-sense to e.g. Conitiation codon, intron of respiratory disease-relevant gene e.g. Con RANTES, MCP4, useful for prophylaxis or treating respiratory disease
                                                                                                                                                                            gene e.g. CCR1,
                                                                                                                                                                       e.g.
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WPI; 2004-293804/27.

Claim 2; SEQ ID NO 1802; 174pp; English.

CC chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)

CC -5 receptor, CCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM,

CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention

CC also relates to a method of screening a candidate compound that binds to

CC one or more nucleic acid target(s) or expressed product(s), for the

CC prevention and/or treatment of a respiratory or lung disease. The

CC oligonucleotides are useful for reducing or inhibiting expression of a

CC gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

CC CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a

CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are

CC useful for preventing or treating a respiratory or lung disease. The

CC useful for preventing or treating a respiratory or lung disease to

CC and/or increased levels of, adenosine and/or levels of adenosine A

CC receptor(s), and/or asthma and/or lung allergies associated with

CC inflammation or an inflammatory disease. The respiratory or lung disease

CC cystic fibrosis (CF), chronic obstructive pulmonary disease (CDPD), The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction o pronchoconstriction. This sequence represents an oligonucleotide of the õ

Sequence 20 BP; 3 A; 7 C; 4 G; 6 T; 0 U; 0 Other;

Matches Query Match Best Local Local 18; Conserv Conservative 1.7%; 0, Score 16.8; Pred. No. 1. Mismatches 1.6e+03 DB 1; 2 Length 20; Indels 0, Gaps 0

밁 Ş 703 AGTTATTCTCCTGCCCCAGC 722 μ AGTGATTCTCCTGCCTCAGC 20

RESULT 1541 Human oligonucleotide #1828. ADO46462 standard; DNA; 20 BP AD046462; (first entry)

Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Estaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; CODD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction. 25-JUL-2003; 2003US-00627930. 11-MAR-2004 US2004049022-A1

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ADO46443
ADO46444
XX
AC ADO4644
AC ADO4644
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AC ADO4644
XX
BT 15-JUL-
XX
DE Human c
XX
KW CCR1; C
KW CCR1; C
KW tryptas
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23-APR-2002; 2002WO-US013143.
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 Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease;
                                                                                                                                                                                                                                                                                                                                                             Sequence 20
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                                                              Human oligonucleotide #1809.
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TANG L.
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ong H;
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The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region codon, coding region, 5' or 3' intron-exon junction, intron or region codon, coding region, 5' or 3' intron-exon junction, intron or region codon, coding region, 5' or 3' end of a nucleic acid target codon codon from a gene encoding interleukin (II) 4 receptor, interleukin (II) 5 receptor, CCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the CC prevention and/or treatment of a respiratory or lung disease. The CC prevention and/or treatment of a respiratory or lung disease a, CC cryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease are cuseful for preventing or treating a respiratory or lung disease. The CC respiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A CC receptor(s), and/or asthma and/or lung allergies associated with CC inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, allergic rhinitis, acute respiratory distress syndrome, pulmonary disease (CCPD), chronic obstructive pulmonary disease (CCPD).
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Shahabuddin S, Lu H,
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lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
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hypertension, lung 1
bronchoconstriction.
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CONG H.
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2002WO-US013143.
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This sequence represents an oligonucleotide
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Matches

18;

Conservative

0;

Mismatches ŏ.

0,

Gaps

0

..6e+03 DB 1;

Length

Similarity

90.0%;

Score 16.8; Pred.

179

AGTAGAGATGGAGTTTCTCC 198

Sequence

20 BP;

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3 C;

7

G; 5 T;

0

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RESULT 1543
ADO45256
The invention relates to oligonucleotides anti-sense to an initiation CC codon, coding region, 5' or 3' intron-exon junction, intron or region CC with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target CC chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-5 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PD24 A, PD24 B, PD24 C or PD24 D. The invention CC also relates to a method of screening a candidate compound that binds to CC one or more nucleic acid target(s) or expressed product(s), for the CC prevention and/or treatment of a respiratory or lung disease. The CC gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PD24 A, PD24 B, PD24 C, or PD24 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The CC respiratory or lung disease is associated with hyper-responsiveness to C reprivation or an inflammatory disease. The respiratory or lung disease.

CC inflammation or an inflammatory disease. The respiratory or lung disease.
                                                                                                                                                                                                                                                                                                                                                                                Novel single or multiple target oligonucleotide anti-sense to e.g. Conitiation codon, intron of respiratory disease-relevant gene e.g. CONNITES, MCP4, useful for prophylaxis or treating respiratory disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Shahabuddin
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(SAND/)
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(AGUI/)
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SHAHABUDDIN :
LU H.
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2002WO-US013143.
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RESULT 1544
AD04526
ID AD04526
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Best Local
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Shahabuddin S, Lu H,
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                                                                                                                       Claim 2;
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(TANG/)
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TANG L.
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ng H;
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The invention relates to oligonucleotides anti-sense to an init codon, coding region, 5' or 3' intron-exon junction, intron or with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid

an initiation acid target

CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention country treatment of a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The coligonucleotides are useful for reducing or inhibiting expression of a gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, cCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, ctryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The cc useful for preventing or treating a respiratory or lung disease. The cc respiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A cc receptor(s), and/or asthma and/or lung allergies associated with cc inflammation or an inflammatory disease. The respiratory or lung disease constructory or lung cryptases (CPD), chronic obstructive pulmonary disease (CPDD), consisted with a constructive pulmonary disease (CPDD).

allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of

of the

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RESULT 1545
AD047047
ID AD04704
XX AD04704
XX AD04704
XX Human;
KW CCR1; C
XX Human;
KW CCR1; C
KW Lryptas
KW Lryptas
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Best Local (
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     Мусе ЈW,
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     Sandrasagra A,
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2002WO-US013143.
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     Aguilar D,
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Codon, coding region, 5' or 3' intron-exon junction, intron or region codon, coding region, 5' or 3' intron-exon junction, intron or region comes from a gene encoding interleukin (II) 4 receptor, interleukin (II) 6-5 receptor, CCR1, ECCR3, 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region
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                                                                                                allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide o
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밁 S Matches Query Match Best Local Similarity 864 GCTGGGATTACAGGCGTGAG 883 1 GCTGGGATTATAGGCATGAG 20 18; Conservative 90.0%; 0; Mismatches Score 16.8; Pred. No. 1.6e+03 DB 1; 2 Length Indels 0, Gaps 0

Sequence 20

BP; 5

A; 2 C; 8

G; 5 T; 0 U; 0 Other;

Human oligonucleotide ADO45254 standard; DNA; 15-JUL-2004 ADO45254; (first entry) #620. 20 ВP

Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PD84 A; PD84 B; PD84 C; PD84 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction. US2004049022-A1 Homo sapiens (년 (년

11-MAR-2004

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Best Local Similarity
Matches 18; Conser
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Shahabuddin S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       bronchoconstriction. This sequence represents an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          is Chosen from airway inflammation, allergy, asthma, impeded respiration, systic fibrosis (CF), chronic obstructive pulmonary disease (COPD),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NYCE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SHAH/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WILL/)
                                                                                                                                                                                                                                                                                                                                                            537 CCTGCCTCAGCCTCCCAAGT 556
                                                                                                                                                                                                                                                                                                                                                                                                       18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ) MILLER S.
) SHAHABUDDIN S.
) LU H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TANG L.
AGUILAR D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NYCE J W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CONG H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SANDRASAGRA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 620;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sandrasagra A, Tang L,
in S, Lu H, Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 2 A;
                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                         (first entry)
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2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                               1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                            20
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                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                               Score 16.8;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Miller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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Chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-6 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to CC one or more nucleic acid target(s) or expressed product(s), for the CC prevention and/or treatment of a respiratory or lung disease. The CC oligonucleotides are useful for reducing or inhibiting expression of a CC gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CC CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC useful for preventing or treating a respiratory or lung disease. The CC respiratory or lung disease is associated with hyper-responsiveness to CC and/or increased levels of, adenosine and/or levels of adenosine A CC receptor(s), and/or asthma and/or lung allergies associated with CC inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, CC cystic fibrosis (CP), chronic obstructive pulmonary disease (COPD),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target with 2-10 nucleotides of the interleukin (IL)-4 receptor, interleukin (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shahabuddin S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nyce JW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SAND/)
(TANG/)
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allergic rhinitis, acute respiratory distress syndrome, pulmonary
hypertension, lung inflammation, bronchitis, airway obstruction o
bronchoconstriction. This sequence represents an oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; SEQ ID NO 633; 174pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-293804/27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MILLER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SANDRASAGRA A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SHAHABUDDIN
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in S, Lu H,
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2002WO-US013143
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Cong ;
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ong H;
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       of the
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Query Match Best Local S Matches 18

Local Similarity

90.0%;

Score 16.8; D Pred. No. 1.6e 0; Mismatches

.6e+03

Length 20; Indels

0

Gaps

0

0,

Sequence

20

BP;

4 A;

œ C;

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2 T; 0

U; 0 Other;

Human; 88; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5;

Human oligonucleotide #633

61 0 140

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ADO45255 standard;
                                     GGACTACAGGCGCCCGCTAC
 DNA;
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20

lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; (chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; I CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; ling disease; humanisms. Human oligonucleotide #621 15-JUL-2004 (first entry bronchoconstriction

Ç

US2004049022-A1 Homo sapiens

11-MAR-2004

23-APR-2002; 23-APR-2002; 25-JUL-2003; 2003US-00627930. 2002WO-US013135. 2002WO-US013143.

NYCE J W

(MILL/) (TANG/) (/HAHS AGUI/) CONG H. TANG SANDRASAGRA SHAHABUDDIN MILLER S. AGUILAR D. ×

Shahabuddin Ä Sandrasagra in S, Lu H, · Cong Tang L, ong H; Aguilar D, Miller S

WPI; 2004-293804/27

Novel single or multiple target oligonucleotide anti-sense to e.g. Conitiation codon, intron of respiratory disease-relevant gene e.g. Corantes, MCP4, useful for prophylaxis or treating respiratory disease e.g. CCR1, e.g

Claim 2; SEQ ID NO 621; 174pp; English

RESULT 1548
AD045255
AD045255
AC AD04525
AC and/or asthma and/or lung allergies associated with

codon,

coding region,

5 to

oligonucleotides a or 3' intron-exon

anti-sense n junction,

to an initiation intron or region

The invention relates

2

SEQ

ij

NO 725;

174pp;

English

Novel single or multiple target oligonucleotide anti-sense to e.g. Conitiation codon, intron of respiratory disease-relevant gene e.g. Convers, MCP4, useful for prophylaxis or treating respiratory disease

CCR1,

e.g.

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RESULT 1549
ADO45359
ID ADO4535
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-1 CCR1; CCR3; Estaxin-1; RAWTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        is chosen from airway inflammation, allergy, asthma, impeded resp. cystic fibrosis (CF), chronic obstructive pulmonary disease (COPD), allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the characteristic of the control of the cont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 3
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Shahabuddin
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                                                                                                                                                                                                         WPI; 2004-293804/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                             (NYCE/)
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23-APR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                            (TANG/)
                                                                                                                                                                                                                                                                                                                    (CONG/)
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                                                                                                                                                                                                                                                                                                                                                         MILLER S.
SHAHABUDDIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NYCE J W.
                                                                                                                                                                                                                                                                                                                                                                                                   AGUILAR D.
                                                                                                                                                                                                                                                                                                                                                                                                                            TANG
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2002WO-US013143.
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Cong
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ong H;
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Pred.
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No. 1
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                                                                                                                                                                                                                                                                              Miller
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disease (COPD),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CF;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC chosen from a gene encoding interleukin (II). 4 receptor, interleukin (II) control of tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The CC oligonuclectides are useful for reducing or inhibiting expression of a cegene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CC CR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a constituting or treating a respiratory or lung disease. The CC useful for preventing or treating a respiratory or lung disease to cuseful for preventing or treating a respiratory or lung disease. The CC cand/or increased levels of, adenosine and/or levels of adenosine A Creceptor(s), and/or asthma and/or lung allergies associated with CC inflammation or an inflammation, allergy, asthma, impeded respiration, collection for allergies respiratory or lung disease. CC cystic fibrosis (CF), chronic obstructive pulmonary disease (COPD),
                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-1 CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airham; lung allergy; inflammation; inflammatory disease; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human oligonucleotide #1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADO46444 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the
                                                                                                    (NYCE/)
(SAND/)
(TANG/)
                                                                                                                                                                                      23-APR-2002;
23-APR-2002;
                                                                                                                                                                                                                                                       25-JUL-2003; 2003US-00627930
                                                                                                                                                                                                                                                                                                                                           US2004049022-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JUL-2004
                                                                             AGUI/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target
                                                                                                                                                                                                                                                                                                                                                                                                                           inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        647 GGCTGGAGTGCAGTGGCGCA 666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    _
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18;
             MILLER S.
SHAHABUDDIN (
                                                                                                                                              NYCE J W
                                                                                                      TANG L.
                                                                                                                        SANDRASAGRA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGCTGGAGTGAAGTGGCACA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                      2002WO-US013135
2002WO-US013143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                         bronchitis; airway obstruction; bronchoconstriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>.</u>.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 16.8; DB 1
Pred. No. 1.6e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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Sequence

20 BP; 3

A; 5 C; 6

G; 6 T; 0 U; 0 Other;

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CC also relates to a method of screening a candidate compound that binds to come or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The collyonucleotides are useful for reducing or inhibiting expression of a gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CC (CR1, CCR3, Echavin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The CC useful for preventing or treating a respiratory or lung disease. The CC respiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A creceptor(s), and/or asthma and/or lung allergies associated with CC inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, CC ystic fibrosis (CP), chronic obstructive pulmonary disease (CPD), cronic obstruction or hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-entron-3'-end of a nucleic acid target chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL-5 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel single or multiple target oligonucleotide anti-sense to e.g. Conitiation codon, intron of respiratory disease-relevant gene e.g. Corners, MCP4, useful for prophylaxis or treating respiratory disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nyce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; SEQ ID NO 1811; 174pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ä,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sandrasagra
in S, Lu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    · A, Cong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tang L,
ong H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Miller
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RESULT 1551
ADO46437
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                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local
                                                                               Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Botaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; inflammation; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension;
                                                                                                                                                                                                                Human oligonucleotide #1803.
                                                                                                                                                                                                                                                                                                 ADO46437 standard; DNA;
                                         Homo sapiens
                                                                                                                                                                                                                                          15-JUL-2004
                                                                     inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                  791 GGGGTTCACCATGTTCGCCA 810
                                                                                                                                                                                                                                                                                                                                                                                                                              18;
                                                                                                                                                                                                                                                                                                                                                                         _
                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                        GGGTTTCACCATGTTGGCCA 20
                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                          1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 16.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                            1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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RESULT 1552
AD011743/c
ID AD01174
XX
AC AD01174
AC AD01174
XX
DT 15-JUL-
XX
DT 15-JUL-
XX
DE Single
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                                                                                                                                                                                      밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL)-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The CC oligonucleotides are useful for reducing or inhibiting expression of a gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, Ctryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The CC useful for preventing or treating a respiratory or lung disease. The creeptor(s), and/or asthma and/or lung allergies associated with CC inflammation or an inflammatory disease. The respiratory or lung disease inflammation or an inflammatory disease. The respiratory or lung disease incled respiration.
                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-APR-2002; 2002WO-US013135.
23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-JUL-2003; 2003US-00627930
                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                          allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-293804/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nyce JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (AGUI/)
                                                                                                                                                                                                                                                                                                                                                                          bronchoconstriction.
   Single multiplex PCR
                                                                                                       ADO11743 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                           is chosen from airway inflammation, allergy, asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary disease (COPD),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (TANG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SAND/)
                                    15-JUL-2004
                                                                      ADO11743;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SHAH/)
                                                                                                                                                                                                                      1023 CTCCCAAGCAGCTGGGATTA 1042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ) NYCE J W.
) SANDRASAGRA A.
) TANG L.
                                                                                                                                                                                                                                                        18;
                                                                                                                                                                                      _
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CONG H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LU H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MILLER S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SHAHABUDDIN S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AGUILAR D.
                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 1804; 174pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sandrasagra A, Ti
in S, Lu H, Cong
                                                                                                                                                                                      CTCCCCAGTAGCTGGGATTA 20
                                                                                                                                                                                                                                                        Conservative
                                      (first entry)
                                                                                                       DNA;
                                                                                                                                                                                                                                                                      1.7%;
                                                                                                                                                                                                                                                                                                                         6 C; 5
                                                                                                                                                                                                                                                                                                                                                                          This sequence
   primer #1115.
                                                                                                       20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tang L,
ong H;
                                                                                                       ВP
                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                         G; 5 T; 0
                                                                                                                                                                                                                                                                       Pred.
                                                                                                                                                                                                                                                                                       Score 16.8;
                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                         No.
                                                                                                                                                                                                                                                                                                                                                                          represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Aguilar D,
                                                                                                                                                                                                                                                                                                                         U; 0 Other;
                                                                                                                                                                                                                                                                         .6e+03
                                                                                                                                                                                                                                                                                       DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                          an oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Miller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ŝ
                                                                                                                                                                                                                                                         <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                             S
S
                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                         0
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문 S

0

RESULT 1553

ADO52208 standard; DNA; 20

ВP

**EXEXEXEX** 

12-AUG-2004 AD052208

(first

entry)

cytostatic; gene therapy; inhibitors of apoptosis-like; IAP-like; Human inhibitor of apoptosis-like antisense oligonucleotide seqid 82

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CC chain reaction by aligning a first primer and a second primer. The method comprises: (a) aligning a first primer and a second primer; and (b) contain four or more bases that are perfectly matching to the 3' end does not contain four or more bases that are perfectly matching to the 3' end cost and one so the first primer or a second primer, the first primer at its cases that are perfectly matching to the 3' end cost and does not contain seven or more bases that are perfectly matching cases that are perfectly matching cases that are perfectly matching to the first primer or the second primer, the first primer at its 3' end does not contain six or more bases that are perfectly matching to a sequence of the first primer or the first primer at its 3' end does not contain six or more bases that are perfectly matching except come mismatch to a sequence anywhere of the first primer at its 3' end cost not contain eleven or more bases that are perfectly matching except come mismatch to a sequence anywhere of the first primer or the second come mismatch to a sequence anywhere of the first primer or the second come mismatch to a sequence anywhere of the first primer or the second come mismatch to a sequence anywhere of the first primer or the second come mismatch to a sequence anywhere of the first primer or the second come mismatch to a sequence anywhere of the first primer or the second come mismatch to a sequence anywhere of the first primer or the second come mismatch to a sequence anywhere of the first primer at its 3' end does not contain second come mismatch to a sequence of the first primer at its 3' end come mismatch to a sequence of the first primer at its 3' end come mismatch to a sequence of the first primer at its 3' end come mismatch to a sequence of the first primer at its 3' end come mismatch to a sequence of the first primer at its 3' end come mismatch to a sequence of the first primer or the second come mismatch to a sequence of the first primer at its 3' end does not contain second come mismatch to a 
                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ss; primer; simultaneous amplification; single multiplex polymerase chain reaction; multifactorial disease; genetic alteration; pharmacogenetic reaction; genotyping; polymorphism; gene expression profiling.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase chain reaction, for high throughput multiplex DNA sequence amplification, comprises aligning two primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a method of designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase chain reaction by aligning a first primer and a second primer. The method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-OCT-2002; 2002US-0417009P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-OCT-2003; 2003WO-US031874
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 38; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-340914/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 6 A; 7 C; 5 G; 2 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          alterations, the studies in pharmacogenetic reactions, the genoty
genetic polymorphisms in a large population, the gene expression
profiling in various samples and high throughput genotyping tech
This sequence corresponds to an example of a primer of the inven
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYNE-) UNIV NEW JERSEY MEDICINE & DENTISTRY.
                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence corresponds
                                                                                                    636 TCTGTCACCCAGGCTGGAGT 655
20
                                                                                                                                                                                                   18;
                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                       1.7%;
                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                       Score 16.8; DB 1;
Pred. No. 1.6e+03;
                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                0 Other;
                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                             2:
                                                                                                                                                                                                                                                                                                       Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the genotyping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              technologies
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RESULT 1554
ADO52272/c
ID ADO52272 standard; DNA; 20
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                                                                                                                                                                                                         The invention describes a compound 8-80 nucleobases in length targeted to CC a nucleic acid molecule encoding inhibitors of apoptosis (IAP)-like, where the compound specifically hybridises with the nucleic acid molecule cc encoding IAP-like comprising 16000 bp (SEQ ID NO. 4) and inhibits the CC expression of IAP-like. Also described are: inhibiting the expression of IAP-like, a screening for a modulator of IAP-like; a CC diagnostic method for identifying a disease state comprising identifying the presence of IAP-like in a sample using at least one of the primers CC selected from 2 sequences comprising SEQ ID NO. 5 or 6, or the probe CC comprising SEQ ID NO. 7; a kit or assay device comprising the compound; and treating an animal having a disease or condition associated with IAP-like. The compound is useful for modulating the expression of IAP-like. CC It is also useful for diagnosing or treating diseases associated with CC expression of IAP-like, e.g. a hyperproliferative disorder. This sequence collegnations is a human inhibitor of apoptosis (IAP)-like antisense
                                                                                                                                            Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New compound targeted to a nucleic acid molecule encoding inhibitors apoptosis (IAP)-like and inhibits expression of IAP-like, useful for modulating the expression of IAP-like or for treating, e.g. hyperproliferative disorder.
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IAP-like modulator; IAP-like associated disorder;
hyperproliferative disorder; human; antisense oligonucleotide;
antisense technology; ss.
                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 14; SEQ ID NO 82; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-399725/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-NOV-2002; 2002US-00303325
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                                                                                                                                                          Similarity
                                                                                                              GCCTCCCGGCTCAAGCGAT 1008
                                                                                                                                                                                                        BP; 4 A; 7 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                             Conservative
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/mod_ba
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/mod_base= OTHER
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/note= "OTHER= Phosphorothioate backbone.
are 5-methylcytidines"
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                                                                                                                                                          1.7%;
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                                                                                                                                                          Score 16.8;
Pred. No. 1.
                                                                                                                                             Mismatches
                                                                                                                                                            .6e+03
                                                                                                                                                                            DB 1;
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                                                                                                                                             Indels
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                                                                                                                                            Gaps
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BP.

Query Match

Local Similarity mes 18; Conserv

Conservative

90.0%; 1.7%;

Score 16.8; DB 1; Length 20; Pred. No. 1.6e+03; O; Mismatches 2; Indels

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The invention describes a compound 8-80 nucleobases in length targeted to CC a nucleic acid molecule encoding inhibitors of apoptosis (IAP)-like, CC where the compound specifically hybridises with the nucleic acid molecule encoding IAP-like comprising 16000 bp (SEQ ID NO. 4) and inhibits the expression of IAP-like. Also described are: inhibiting the expression of CC expression of IAP-like, Also described are: inhibiting the expression of CC expression of IAP-like, also described are: inhibiting the expression of CC expression of IAP-like in a sample using for a modulator of IAP-like; a CC diagnostic method for identifying a disease state comprising identifying CC the presence of IAP-like in a sample using at least one of the primers CC comprising SEQ ID NO. 7; a kit or assay device comprising the compound; CC and treating an animal having a disease or condition associated with IAP-CC like. The compound is useful for modulating the expression of IAP-like, c.g. a hyperproliferative disorder. This sequence corrected with an animal inhibitor of apoptosis (IAP)-like antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New compound targeted to a nucleic acid molecule encoding inhibitors apoptosis (IAP)-like and inhibits expression of IAP-like, useful for modulating the expression of IAP-like or for treating, e.g. hyperproliferative disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
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Sequence
                                                              oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 14; SEQ ID NO 146; 58pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC.
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B₽;
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/note= "OTHER= Phosphorothioate backbone. All
are 5-methylcytidines"
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note= "OTHER= 2'-O-Methoxyethyl
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== "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
C; 7
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   G; 4 T; 0 U; 0 Other;
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                                             Ś
                                                                                                              Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                         The invention relates to a novel method for identifying a subject at risk of breast cancer comprising detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a subject at risk of breast cancer, for early diagnosis, prevention and treatment of breast cancer, possibly via gene therapy, as well as to analyse and predict a response to a breast cancer treatment and in clinical drug trials. The current sequence is that of an Extend primer (also described as probe) of the invention which was used to genotype human intercellular adhesion molecule ICAM-1/ICAM-4/ICAM-5 gDNA. ICAM-1 (human rhinovirus receptor;BB2;CD54;call surface glycoprotein p3.58) has been mapped to chromosomal position 19p13.3-p13.2, ICAM-4 (Landsteiner-Wiener blood group;IW) has been mapped to chromosomal position 19p13.3-cen and ICAM-5 (Landsteiner-Wiener blood group;IW) has
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Identifying a subject at risk of breast cancer by detecting the presence of polymorphic variations in the ICAM, MAPK10, KIAA0861, NUMA1 or GALE regions which are associated with breast cancer in a nucleic acid sample
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24-JUL-2003; 2003US-0490234P.
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                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 4; Page 83; 289pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-441051/41.
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                                                                                                                                                                                                                                                                                         (telencephalin)
                                                                                                                 Local Similarity
                                                          1085
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                                          TAGAGGCGGGGTTTCACCAT 1104
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      TAGAGACGGGGTTTCACTAT 20
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                                                                                                        1.7%;
nilarity 90.0%;
Conservative
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                                                                                                                                            Score 16.8;
Pred. No. 1
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                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                         chromosomal position
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                                                                                                                    Gaps
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RESULT 1556

Analysis; gene expression; reverse transcription; primer; cDNA;

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                                                                                                         RESULT 1557
AAQ75719
                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADP46278;
                                                                                                                                                                                                                                                                                         The invention relates to a novel method for identifying a subject at ri of breast cancer comprising detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a subject at risk of breast cancer, for early diagnosis, prevention and treatment of breast cancer, possibly via gene therapy, as well as to analyse and predict a response to a breast cancer treatment and in clinical drug trials. The current sequence is that of an Extend primer (also described as probe) the invention which was used to genotype human Rho family guanine-nucleotide exchange factor KIAA0881 gDNA which has been mapped to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying a subject at risk of breast cancer by detecting the presence of polymorphic variations in the ICAM, MAPKIO, KIAAO861, NUMA1 or GALE regions which are associated with breast cancer in a nucleic acid sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     breast cancer; cytostatic; gene therapy; human; ss; primer; PCR;
single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-NOV-2002; 2002US-0429136P
24-JUL-2003; 2003US-0490234P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADP46278 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-NOV-2003; 2003WO-US037948
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-JUN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chromosome 3q27.3; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rho family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Extend primer 59 used to genotype human KIAA0861 polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 6; Page 99; 289pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-441051/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Roth RB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                        Reverse
                                               04-AUG-1995
                                                                                                                                                                                                                                                          Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         from a
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                                                                        AAQ75719
                                                                                              AAQ75719 standard; DNA;
                                                                                                                                                                                                                       Local
                                                                                                                                                                                   673 GCTCACTGCAACCTCTGCCT 692
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                       transcription
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                guanine-nucleotide exchange factor; KIAA0861;
                                                                                                                                                                                                                                                            BP; 5 A; 3 C; 9 G;
                                                                                                                                                                                                                                                                                   position
                                                                                                                                                                                                          Conservative
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                                               (first
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                                               entry)
                                                                                                                                                                                                                       1.7%;
                                                                                                                                                                                                                                                                                    3q27.3.
                       primer used in cDNA analysis technique.
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                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SM.
                                                                                                                                                                                                                       .6e+03
                                                                                                                                                                                                                                                             0 Other;
                                                                                                                                                                                                                                    DB 1;
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RESULT 1558
AAQ75730
ID AAQ7573
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ75730 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21
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by digestion
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                              Disclosure; Page 8; 11pp; Japanese
                                                                                    by digestion with restriction
                                                                                                          Analysis of cDNA and gene expression -
                                                                                                                                                                                                                                                                                                                               16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                    01-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                         JP06303997-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reverse transcription primer used in cDNA analysis technique.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 2 A; 0 C; 2 G; 17 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                            93JP-00112515
                                                                                                                                                                                                                                                                                                                               93JP-00112515
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Pred. No. 1.
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                                                                                                          amplification of mRNA followed
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RESULT 1559
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Best Local Similarity
Matches 18; Conserv
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Best Local (
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                                                                                                                                                                                     A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c)
                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                      electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                    Disclosure; Page 8; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                   by digestion with restriction
                                                                                                                                                                                                                                                                                                                                                  Analysis of cDNA and gene expression - by amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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                                                                                                                          21 BP; 3 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                             TTTTTATTTTTTTTTAAT 614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21 BP; 2 A; 1 C; 0 G; 18 T; 0 U; 0 Other;
TTTTTTTTTTTTTTAAT 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          restriction enzyme;
                                                             Conservative
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Pred. No. 1.6e+03;
                                                                           Score 16.8; DB 1;
Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                     enzymes.
                                                              Mismatches
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RESULT 1561
AAQ7572
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XX AAQ7572
AC AAQ7572
XX Reverse
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Analysis of cDNA and gene expression - by digestion with restriction enzymes.
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                                                                                                          aggregate;
                                                                                                                                         Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                               04-AUG-1995
                                                                                                                                                                                     AAQ75722;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-APR-1993;
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                                                                                                                     Analysis;
   16-APR-1993;
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                                                                  JP06303997-A
                      16-APR-1993;
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                                                                                                                                                                                                          standard; DNA;
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; restriction enzyme; ss.
                                                                                                                                                                                                                                                                  TTTTTTTTTTTTTTAAT 20
                                                                                                                     gene
                                                                                                           restriction enzyme;
                                                                                                                                                                                                                                                                             TTTATTTTTATTTTTAAT 614
                                                                                                                                                                                                                                                                                                                                                      BP; 2 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page
                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                                                 (first entry)
   . 93JP-00112515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEGRAPH & TELEPHONE
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                      93JP-00112515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8; 11pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
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                                                                                                                                                                                                                                                                                                                     1.7%;
                                                                                                                                                                                                                                                                                                                                                      0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ₽₽
                                                                                                                     reverse transcription;
                                                                                                                                                                                                           ₽₽
                                                                                                                                                                                                                                                                                                            0,
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                                                                                                                                                                                                                                                                                                                      Score 16.8;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                         1.6e+03;
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                                                                                                                     primer; cDNA;
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                           Matches
                                           Query Match
Best Local (
                                                                                                         A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type (labelled reverse transcription primers (GENESEO files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                            WPI; 1995-018287/03
                                                                                  Sequence
                                                                                                                                                                                                                                        Disclosure; Page 8; 11pp; Japanese.
                                                                                                                                                                                                                                                                  Analysis of cDNA and gene
by digestion with restrict
                                                                                                                                                                                                                                                                                                                                       (NITE )
 429 TTTATTTTATTTTTTTAAG 448
                           18;
                                                                                                                                                                                                                                                                                                                                     NIPPON TELEGRAPH
                                           Similarity
                                                                                  21
                                                                                BP; 2 A; 1 C; 1
                            Conservative
                                                                                                                                                                                                                                                                     restriction
                                         1.7%;
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                                                                                                                                                                                                                                                                                  expression
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                                                                                   <u>.</u>
                            <u>,</u>
                                           Score 16.8;
Pred. No. 1
                                                                                                                                                                                                                                                                     enzymes.
                                                                                   17 T; 0 U; 0 Other;
                              Mismatches
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                                            6e+03;
                                                         DB 1;
                                                                                                                                                                                                                                                                                   amplification of mRNA followed
                               Indels
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RESULT 1562
AAQ75712 standard; DNA;
 21
 ВP
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04-AUG-1995 AAQ75712; 01-NOV-1994. Synthetic. aggregate; Analysis; gene Reverse transcription primer used in cDNA analysis technique. JP06303997-A restriction (first entry) expression; enzyme; reverse yme; ss. transcription; primer; CDNA;

16-APR-1993; 93JP-00112515

16-APR-1993; 93JP-00112515

(NITE ) NIPPON TELEGRAPH & TELEPHONE CORP

WPI; 1995-018287/03

Analysis of cDNA and gene expression by digestion with restriction enzymes enzymes. γď amplification of, mRNA followed

Disclosure; Page 7; 11pp; Japanese.

A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. Th method can be used to analyse gene expression rapidly and easily an aggregate plural type <u>۾</u> ۾

Sequence 21 BP; N Þ 0 ç \_ ଦ 18 Ŧ, 0 U; 0 Other;

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RESULT 1563
AAQ75721
ID AAQ75722
XX AAQ7572
AC AAQ7572
XX AAQ7572

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                                                                                                                                                                        RESULT 1564
AAA96626
                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 18; Conserv
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HPRT; enhanced homologous recombination; EHR; recombinase;
gene targeting; gene recombination; phenotype screening; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                 A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in separate lames. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Analysis; gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Reverse transcription primer used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-AUG-1995
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                                                                                        08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 8; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Analysis of cDNA and gene expression -
by digestion with restriction enzymes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-APR-1993;
                                                    PCR primer used to generate a biotinylated 318 bp HPRT probe
                                                                                                                                                          AAA96626 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                    be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                    BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
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Pred. No. 1.
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amplification of mRNA followed
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                                                                                                                                                                                                                                                                                                                 Indels
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   88
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     RESULT 1565
AAZ72283/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  drug targets
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                                                     21-APR-1998;
23-NOV-1998;
                                                                                                                                                                                                                                               diagnosis; ss.
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                                                                                                                                            28-OCT-1999
                                                                                                                                                                          WO9954500-A2
                                                                                                                                                                                                            Homo sapiens.
                     (GEST ) GENSET
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PCR primers AAA96626-27 were used to generate a probe for HPRT. The probis used in the course of the invention. The specification describes a method for cloning a target nucleic acid. The method involves providing an enhanced homologous recombination (EHR) composition comprising a recombinase, a targeting polynucleotide, and a separation group. These are then contacted with a target library, from which the target nucleic acid is isolated, using a robotic system. The EHR technique is useful for acid is isolated, using a robotic system. The EHR technique is useful for acid is isolated.
                                                                                                                                                                                                                                                                                                                          Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cloning a target nucleic acid for gene targeting, recombination, phenotype screening and biovalidation of drug targets, involves utilizing enhanced homologous recombination techniques.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Page 50; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-638261/61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human biallelic marker upstream amplification primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ72283 standard;
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98US-0082614P.
98US-0109732P.
                                                                                  99WO-IB000822.
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Pred. No. 1.
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Claim 1; Page 63; 83pp; English.

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ARAH3978
AAH3978
AC SNP Spe
KW Lesch-N
KW Jesch-N
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOs 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the sequence of the se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cohen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-OCT-2000; 2000WO-US028436.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200129262-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNPE; genotyping;
Lesch-Nyhan syndro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP specific lower PCR primer SEQ ID 2582.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH39786 standard;
                                        absence or identity acid sample.
                                                                             New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nu
                                                                                                                                                                                                                                                                                                                                                                                                                                   15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                              (ORCH-) ORCHID BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000-013267/01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            eotide polymorphism; SNP; single nucleotide primer extension;
yping; agammaglobulinaemia; diabetes insipidus; cancer;
syndrome; muscular dystrophy; familial hypercholesterolaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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Pred. No. 1.
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RESULT 1567
ABS60534/c
ID ABS605
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                                       04-DEC-2000;
23-JAN-2001;
02-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Aminopeptidase P; XPNEP2; bradykinin receptor B1; ds; BDKRB1; tachykinin receptor B1; TACR1; C1 esterase inhibitor; C1NH; kallikrein KLK1; bradykinin receptor B2; BDKRB2; gene therapy; ALK1; bradykinin receptor B2; BCKRB2; grotease inhibitor 4; P14; angiotensin converting enzyme 2; ACE2; protease inhibitor 4; P14; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 3 A; 8 C; 4 G; 6 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  paternity analysis. The present sequence represents for a human SNP containing DNA sequence
                                                                                                                                                                                                                                                                                                                                                                       cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; aneurysm; embolism; thrombosis; coronary artery disease; angioedaema;
                                                                                                                                                                                                         WO200261131-A2
                                                                                                                                                                                                                                                                                                              arteriosclerosis; atherosclerosis; hypersensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human polymorphism associated DNA sequence #283.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABS60534;
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                                                                                                                          03-DEC-2001; 2001WO-US047235.
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(BRIM ) BRISTOL-MYERS SQUIBB
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18; Conserv
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                                     ; 2000US-0251015P.
; 2001US-0263678P.
; 2001US-0273037P.
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Pred. No. 1
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ABS60764/c
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AC ABS607
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DT 05-NOV
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DE Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC 2 (ACE2) or protease inhibitor 4 (PHA), comprising at least one composition. Also included are (1) a probe that hybridises to a composition position as provided in the detailed summary of single conclectide polymorphisms comprising additional 5 and 3 flanking genomic comprising the sample from one or more nucleic acid sample comprising cobtaining the sample from one or more individuals and determining the concleic acid sequence at one or more individuals and determining the concleic acid sequence at one or more polymorphic positions in a gene concleic acid sequence at one or more polymorphic positions in a gene comprising protein selected from the group above; (3) constructing (M2) constructing (M2) an individual at risk of developing a disorder comprises one or more polymorphic positions within a gene encoding a comprise so one or more polymorphic positions within a gene encoding a comprise so one or more polymorphic positions within a gene encoding a comprise so one or more polymorphic positions within a gene encoding a comprise so one or more polymorphic positions within a gene encoding a condition of an ACE inhibitor and/or vasopeptidase which the protein sequence in the specification. Antibodies and biological samples. The present sequence is 
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                                                                                                                                                                                                    RESULT 1568
                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 18
                    Human polymorphism associated DNA sequence #401
                                                                   05-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 801; 977pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-619265/66.
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                                                                                                                 ABS60764;
                                                                                                                                                           ABS60764 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       encoding aminopeptidase P (XPNE tachykinin receptor B1 (TACR1),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           autoimmune diseases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated nucleic acid with at least one polymorphic position, us for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (KLK1), bradykinin receptor B2 (BDKRB2), angiotensin converting enzyme
                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                                                 1085 TAGAGGCGGGGTTTCACCAT 1104
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                                                                   (first entry)
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                                                                                                                                                           ВP
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                                                                                                                                                                                                                                                                                                                                                                               Score 16.8; DB 1;
Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
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Aminopeptidase P; XPNEP2; bradykinin receptor B1; ds; BDKRB1; tachykinin receptor B1; TACR1; C1 esterase inhibitor; C1NH; kallikrein 1; KIKI; bradykinin receptor B2; BDKRB2; gene therapy; angiotensin converting enzyme 2; ACE2; protease inhibitor 4; P14; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma; cardiovascular disease; angina pectoriis; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; angioedaema; aneurysm; embolism; thrombosis; coronary artery disease; angioedaema; arteriosclerosis; atherosclerosis; hypertensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD; Chronic obstructive pulmonary disease; enterocolitis.

W0200261131-A2.

03-DEC-2001; 2001WO-US047235.

04-DEC-2001; 2001WO-US047235.

04-DEC-2001; 2001WO-US047235.

05-MAR-2001; 2001US-0273037P.

(BRIM) BRISTOL-MYERS SQUIBB CO.

(TSUC/) TSUCHIHASHI Z.
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Tsuchihashi Z, Hui L, Zerba KE, Ma-Edmonds M, Perrone MH. Swanson BN, Powell JR;

WPI; 2002-619265/66.

New isolated nucleic acid with at least one polymorphic position, useful for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and autoimmune diseases.

Disclosure; Page 876; 977pp; English

CC encoding aminopeptidase P (XPMEP2), bradykinin receptor B1 (BDKRB1), CC tachykinin receptor B1 (TACR1), C1 esterase inhibitor (CINH), kallikrein CC 1 (KLK1), bradykinin receptor B2 (BDKRB1), angiotensin converting enzyme C2 (ACE2) or protease inhibitor 4 (P14), comprising at least one CC polymorphic position. Also included are (1) a probe that hybridises to a C polymorphic position as provided in the detailed summary of single cultivated in the detailed summary of single complete cide polymorphisms comprising additional 5, and 3, flanking genomic cobtaining the sample from one or more polymorphic positions in a gene ce encoding a protein selected from the group above; (3) constructing (M2) to dientifying (M3) an individual at risk of developing at dientifying (M3) an individual at risk of developing a disorder culsing the polymorphic data; (5) a library of mucleic acids each separate polymorphic positions in a gene encoding a protein selected from the group above; (3) constructing (M2) c human protein selected from the group above; (3) constructing (M2) c human protein selected from the group above; and (6) genetyping at least to polymorphic data; (5) a library of mucleic acids, each of which comprises one or more polymorphic positions within a gene encoding a nucleic acid sample, determining the nucleicide present in at least one polymorphic position, and comparing at C and composition with a known data set. The genes, (M1, M2, M3 and M4) c and composition with a known data set. The genes, (M1, M2, M3 and M4) c preventing various disorders such as angioedaema and diseases which c rivolve angiogenesis like haemangiomas, tunours, sarcomas, Crohn's c disease, trachomas, and cardiovascular diseases like angina pectoris, c hypertension, heart failure, myocardial infarction, ventricular constructive pulmonary diseases (COPD) and entercolitis (many other constructive pulmonary diseases (COPD) and entercolitis (many other constructive pulmonary disease (COPD) and entercolitis (many other constructive pulmonary dise

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ABS60765/c
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DE Human
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Aminog
KW Aminog
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KW Amicoi
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Acteri
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Best Local
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23-JAN-2001;
02-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABS60765 standard; DNA;
                                                                                                                                                                                                                                                                                        New isolated nucleic acid with at least one polymorphic position, useful for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tsuchihashi
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      1 (\dot{\text{KLK1}}), bradykinin receptor B2 (BDKRB2), angiotensin converting enzyme 2 (ACE2) or protease inhibitor 4 (PI4), comprising at least one polymorphic position. Also included are (1) a probe that hybridises to a
                                                                                             The invention relates to an isolated nucleic acid from a human encoding aminopeptidase P (XPNEP2), bradykinin receptor B1 (BDI tachykinin receptor B1 (TACR1), C1 esterase inhibitor (C1NH), )
                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-619265/66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BRIM )
                                                                                       encoding aminopeptidase tachykinin receptor B1
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TSUCHIHASHI Z.
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Powell JR;
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; 2001US-0263678P.
; 2001US-0273037P.
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                                                                                                                                                                                                          Page 876;
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                                                                                                                                                                                                       977pp; English.
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                                                                                                                           (BDKRB1)
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                                                                                                kallikrein
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cc polymorphic position as provided in the detailed summary of single cnucleotide polymorphisms comprising additional 5' and 3' flanking genomic cc gequence; (2) analysing (M1) at least one nucleic acid sample comprising controlled acid sample comprising controlled acid sample comprising controlled acid sample comprising controlled acid sample comprising the sample from one or more individuals and determining the comprising constructing (M2) constructing (M3) an individual at risk of developing a disorder cupon administration of an ACE inhibitor and/or vasopeptidase inhibitor constructing the polymorphic data; (5) a library of nucleic acids, each of which comparises one or more polymorphic positions and (6) genotyping (M4) and construction protein selected from the group above; and (6) genotyping (M4) and construction with a known data set. The genes, (M1, M2, M3 and M4) construction with a known data set. The genes, (M1, M2, M3 and M4) constructions are useful for detecting, diagnosing, treating, construction was an angioedaema and diseases which constructs and cardiovascular diseases like angina pectoris, construction, wentricular convertions, heart failure, myocardial infarction, ventricular convertions, ventricular diseases, inflammatory constructive pulmonary disease, aneurysm, embolism, thrombosis, coronary constructive pulmonary disease, sepsis, autoimmune diseases, inflammatory constructive pulmonary disease (COPI) and enterocolitis (many other construction but is not referred to anywhere else in the specification. Antibodies and biological samples. The present sequence is included in the sequence construction of t
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ABS60535/c
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Best Local S
Matches 18
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Pred. No. 1
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ABS60535;
                                ABS60535 standard; DNA;
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05-NOV-2002

(first

entry)

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Aminopeptidase P; XPNEP2; bradykinin receptor B1; ds; BDKRB1; tachykinin receptor B1; TACR1; C1 esterase inhibitor; C1NH; kallikreii KuK1; bradykinin receptor B2; BDKRB2; gene therapy; angiotensin converting enzyme 2; ACE2; protease inhibitor 4; P14; polymorphism; haemangiona; tumour; sarcoma; Crohn's disease; trachoma, cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; angioedaema; arteriosclerosis; thrombosis; coronary artery disease; angioedaema; arteriosclerosis; atherosclerosis; hypersensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD; Chronic Chessia; and coronary arthritis; cancer; wound; chessia; cancer; cancer; cancer; cancer; cancer; cancer; cancer; canc
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Homo sapiens.
                                                                                                                                                                                                                                           pulmonary
                                                                                                                                                                                                                                                       disease;
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03-DEC-2001; 2001WO-US047235

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RESULT 1571
ABQ93617
ID ABQ9361
XX
AC ABQ9361
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Best Local Sim:
Matches 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           conclaining the sequence at one or more polymorphic positions in a gene encoding a protein selected from the group above; (3) constructing (M2) chaplotypes using the genes comprising grouping at least two nucleic acids (4) identifying (M3) an individual at risk of developing a disorder using the polymorphic data; (5) a library of nucleic acids; (4) identifying (M3) an individual at risk of developing a disorder using the polymorphic data; (5) a library of nucleic acids, each of which comprises one or more polymorphic positions within a gene encoding a human protein selected from the group above; and (6) genotyping (M4) an individual comprising obtaining a nucleic acid sample, determining the nucleotide present in at least one polymorphic position, and comparing at least one position with a known data set. The genes, (M1, M2, M3 and M4) and compositions with a known data set. The genes, (M1, M2, M3 and M4) and compositions are useful for detecting, diagnosing, treating, at least one position with a known data set. The genes, (M1, M2, M3 and M4) and compositions are useful for detecting, diagnosing, treating, compositions are useful for detecting, diagnosing, treating, involve angiogenesis like haemangiomas, tumours, sarcomas, Crohn's disease, trachomas, and cardiovascular diseases intermining apectoris, hypertrension, heart failure, mycoardial infarction, ventricular hypertrension, heart failure, mycoardial infarction, ventricular hypertrension, heart failure, acousting and hypertrension; arteriosclerosis and/or atherosclerosis, and hypertrension; arteriosclerosis and/or atherosclerosis, and
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23-JAN-2001;
02-MAR-2001;
                                                ABQ93617 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   encoding aminopeptidase P (XPNEP2), bradykinin receptor B1 (BDKRB1), tachykinin receptor B1 (TACR1), C1 esterase inhibitor (C1NH), kallikrein tachykinin receptor B2 (BDKRB2), angiotensin converting enzyme 2 (ACE2) or protease inhibitor 4 (P14), comprising at least one polymorphic position. Also included are (1) a probe that hybridises to a polymorphic position as provided in the detailed summary of single nucleotide polymorphism comprising additional 5' and 3' flanking genomic sequence; (2) analysing (M1) at least one nucleic acid sample comprising obtaining the sample from one or more individuals and determining the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated nucleic acid with at least one polymorphic position, useful for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tsuchihashi Z, Hui L,
Swanson BN, Powell JR;
                                                                                                                                                                                                                                                                                                                                                      Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                   polynucleotides are also useful for chromosome identification. Antibodies against the proteins may be utilised for immunophenotyping of cell lines and biological samples. The present sequence is included in the sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hypersensitivity reactions, sepsis, autoimmune diseases, inflammatc arthritis, cancer, wounds, viral, bacterial or fungal infection, Chobstructive pulmonary disease (COPD) and enterocollitis (many other diseases and disorders are listed in the specification). The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an isolated nucleic acid from a human gene encoding aminopeptidase P (XENEP2), bradykinin receptor B1 (BDKXB1)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                autoimmune diseases.
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                                                                                                                                                                                                                                                                                                                                                         BP; 5
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2001US-0263678P.
2001US-0273037P.
                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                       referred
                                                                                                                                                                                                                                                                               1.7%;
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to anywhere else in
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                                                                                                                                                                                                                                                                                    16.8;
No. 1.
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Best Local (
                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a novel Disrupted-In-Schizophrenia (DISC) 1 allelic variant polynucleotide. The polypetides of the invention have neuroleptic activity. The polynucleotides may have a use in gene therapy. DISC1 or DISC2 nucleic acid molecules are useful for diagnosing or treating a subject having a disease or disorder associated with specific DISC1 or DISC2 alleles and/or aberrant DISC1 expression or activity e.g. neuropsychiatric disorder such as schizoaffective, bipolar, unipolar affective or adolescent conduct disorder or schizophrenia. Similarly, the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New human Disrupted-In-Schizophrenia (DISC) 1 and DISC2 genes containing single nucleotide polymorphisms, useful for preventing or treating neuropsychiatric disorders e.g. schizophrenia.
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                              compound that inhibits DISC1 protein activity may be used in the method for treating such neuropsychiatric disorders. The sequences shown in ABQ993575-ABQ93558 represent the PCR primers used in the invention to amplify the sequences of DISC2 and DISC2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; Disrupted In Schizophrenia 1; DISC1; neuroleptic; gene therapy; neuropsychiatric disorder; schizoaffective disorder; bipolar disorder; unipolar affective disorder; adolescent conduct disorder; schizophrenia;
               WO200227035-A2
                                                                      Enhanced homologous recombination; EHR; recombinase; hybridisation; gene targeting; nucleic acid isolation; HPRT; PCR; primer; ss.
                                                                                                              HPRT probe generating primer hExo3-2A.
                                                                                                                                             30-JUL-2002
                                                                                                                                                                            ABL58478;
                                                                                                                                                                                                      ABL58478 standard; DNA;
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                                         Synthetic
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                                                                                                                                                                                                                                                                                                                                         Conservative
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Pred. No. 1.
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               phenotype screening, biovalidation of drug targets, DNA cloning, DNA modification, isolation of gene families, orthologues and paralogues, identification of alternatively spliced isoforms, gene mapping, disgnostic testing for single and multiple nucleotide polymorphisms, disgnostic testing for single and multiple nucleotide polymorphisms, differential gene expression and genetic profiling, nucleic acid library production, subtraction and normalization, in situ gene targeting (hybridisation) in cells, in situ gene recombination in cells and animals, high throughput phenotype screening of cells and animals, high throughput phenotype screening of cells and animals, and biovalidation of drugs in transgenic recombinant cells and animals. Sequences ABLS8478-79 represent PCR primers for generating and animals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to high throughput integrated genomics, or isolating target nucleic acid (NA) or genomic DNA. The method invol contacting an enhanced homologous recombination (EHR) composition comprising a recombinase, a first and second target polynucleotide complementary to each other and a separation group, with a library or DNA or with one or more NA samples under conditions favouring the polynucleotide is useful for gene targeting, recombinating the polynucleotide is useful for gene targeting, recombinating the polynucleotide is useful for gene targeting, recombinating the polynucleotide is useful for gene targeting.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-APR-2002
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                                                                                                                                                                                                                                                                                                     Human;
renal (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABS66754 standard;
                               26-JAN-2001;
                                                                                   25-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      493
                                                                                                                                                                                                                                                                                                  multidrug resistance-associated protein 1; MRP-1; ss; cancer; cytostatic; single nucleotide polymorphism.
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                                                                                                                                                                                                                                                                                                                                                                                      polymorphic DNA region #19
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                               2001EP-00101651
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Pred. No. 1.6e
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                                                                                                                                                                 -1) polymucleotide. The polymucleotide is useful in an in vitro method for identifying a single nucleotide polymorphism and for identifying and obtaining a pro-drug or drug capable of modulating the activity of a molecular variant of MRP-1 or for identifying and obtaining an inhibitor of the activity of a molecular variant of MRP-1. The sequences are useful for diagnosing a disorder related to the presence of a molecular variant of MRP-1 or susceptibility to such a disorder, where the disorder is cancer (particularly renal cancer) or a disease related to multidrug
                                                                                                                                                                                                                                                                                                                                                                                        Novel
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                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                           Example
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                                                                                                                                                                                                                                                                                                invention relates to
                    834 TGTGATCTGCCTGCCTCGGC 853
                                                                                                                                                                                                                                                                                                                                                      multidrug resistance-associated protein 1 osis and treatment of cancer and multidrug ses, and for identifying single nucleotide
                                                              18;
                                                                                                                                                                                                                                                                                                                           <u>ب</u>
                                                                           Similarity
                                                                                                                         21
                                                                                                                                                                                                                                                                                                                         Page 66; 198pp; English.
                                                                                                                         ₿P;
                                                                                                                                                      This sequence represents a human MRP-1 polymorphic
                                                              Conservative
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                                                                                                                        1 A;
                                                                                                                                                                                                                                                                             The polynucleotide is useful in an in vitro method
                                                                           1.7%;
                                                                                                                         8 C; 7 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                a multidrug
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                                                                                           Score 16.8;
                                                                Mismatches
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                                                                              No.
                                                                                                                                                                                                                                                                                                resistance-associated protein 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                      M
                                                                            1.6e+03
                                                                                              B
                                                                                                                                                                                                                                                                                                                                                          polynucleotide useful resistance related polymorphisms.
                                                                                           Length
                                                                Indels
                                                                                              21;
                                                                0
                                                                                                                                                         DNA region
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RESULT 1574
ABS66755/c
WO200259142-A2
                                  Homo sapiens.
                                                                    Human; multidrug resistance-associated protein 1; MRP-1; ss; renal cancer; cytostatic; single nucleotide polymorphism.
                                                                                                                           Human MRP-1
                                                                                                                                                                 29-NOV-2002
                                                                                                                                                                                                   ABS66755;
                                                                                                                                                                                                                                     ABS66755 standard;
                                                                                                                                                                                                                                                                                                                                TGTGATCGGCCCGCCTCGGC 20
                                                                                                                             polymorphic DNA region #20
                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                       21
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01-AUG-2002

26-JAN-2001; 2001EP-00101651 25-JAN-2002; 2002WO-EP000796

for

polynucleotide. The pidentifying a single aining a pro-drug or c

invention relates to a multidrug resistance-associated protein 1 polynucleotide. The polynucleotide is useful in an in vitro metho

nucleotide polymorphism andrug capable of modulating

in an in vitro and for identia

method and

identifying ctivity of a

Example

2; Page 66; 198pp; English.

Novel multidrug resistance-associated protein 1 polynucleotide useful diagnosis and treatment of cancer and multidrug resistance related diseases, and for identifying single nucleotide polymorphisms.

for

WPI; 2002-657475/70.

Brinkmann

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Hoffmeyer

S

Mornhinweg

(EPID-)

EPIDAUROS BIOTECHNOLOGIES AG.

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ADC42667/c
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          molecular variant of MRP-1 or for identifying and obtaining an inhibitor of the activity of a molecular variant of MRP-1. The sequences are useful for diagnosing a disorder related to the presence of a molecular variant of MRP-1 or susceptibility to such a disorder, where the disorder is cancer (particularly renal cancer) or a disease related to multiduo
                                                                                                                                                              Fanconi Anaemia (FA)/BRCA pathway gene or protein for the presence of a cancer-associated defect, where the presence of one or more cancer-associated defects is indicative of cancer or an increased risk of cancer in the patient. The method of the invention has cytostatic activity. The method is useful for determining if a patient has cancer, or is at
                                             increased risk of developing cancer, e.g. breast, ovarian or prostate cancer. A microarray of the invention is useful for determining if a patient has cancer, or is at increased risk of developing cancer, by hybridising a nucleic acid sample to the nucleic acid sequences from the array, and detecting the presence of mutations in FA/BRCA pathway genes in the nucleic acid sample from the patient, where detecting the presence of mutations is indicative of a patient who has cancer, or is at
                                                                                                                                                                                                                                                                                                        Example 14; Page 103; 160pp; English
                                                                                                                                                                                                                                                                                                                                                       Diagnosing or determining cancer or increased risk of cancer in a patient, by testing Fanconi Anemia/BRCA pathway gene or protein for cancer-associated defect, that indicates cancer or increased risk of
                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-441436/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               D'andrea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-NOV-2001; 2001US-00998027.
02-NOV-2001; 2001WO-US045561.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chemosensitising;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer; Fanconi Anaemia; FA; BRCA; cytostatic; microarray;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human FANCD2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADC42667;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                            patient
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADC42667 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              resistance. This sequence represents a human MRP-1 polymorphic DNA region
increased risk of developing cancer. A method of the invention is for screening a chemosensitising agent, and the agent obtained is for treating a patient having a cancer. The present sequence is us
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYOR-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (DAND )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                          invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18;
                                                                                                                                                                                                                                                     antion relates to a novel method of diagnosing has cancer or is at increased risk of cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DANA FARBER CANCER INST
UNIV OREGON HEALTH SCI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Taniguchi
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Pred. No. 1.
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RESULT 1576
ADC42296
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                              The invention relates to a method of isolating a target nucleic acid or genomic DNA. The method is useful for isolating a target nucleic acid, e.g. a portion of a target gene, a regulatory sequence, or a nucleic acid comprising single nucleotide polymorphism (SNP), a target genomic DNA, e.g. mammalian chromosome which is a fragment of genome separated from cDNA. The method provides high-throughput integrated genomics useful for phenotype screening, isolation of full-length cDNA clones, identification of functional domains, validation of selected sequence, gene targeting, recombination and biovalidation of drug targets. The present sequence represents the hypoxanthine phosphoribosyl transferase HFRT PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ss; primer; PCR; hypoxanthine phosphoribosyl transferase;
nucleic acid isolation; high-throughput integrated genomion
phenotype screening; gene targeting; drug target biovalida
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Sequence
                                                                                                                                                                Example 1;
                                                                                                                                                                                              Isolating a target nucleic acid or genomic DNA comprises using an enhanced homologous recombination composition and contacting with library of target nucleic acid or genomic DNA library using a rob
                                                                                                                                                                                                             enhanced
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                                                                                                                                                                                                                                                                     Zarling
                                                                                                                                                                                                                                                                                            (LEHM/)
                                                                                                                                                                                                                                                                                                                  (ZARL/)
(CASP/)
(STEP/)
(SERG/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hypoxanthine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20
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18; Conserv
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                                                                                                                                                                                                                                                                                            LEHMAN
PATI S.
                                                                                                                                                                                                                                                                                                                                          ZARLING
CASPI R.
                                                                                                                                                                                                                                                                                                                   STEPHENS K M. SERGEANT R G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
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21
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B₽;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         phosphoribosyl transferase,
                                                                                                                                                                                                                                                                     Caspi R,
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Pred. No. 1.
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 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ed genomic;
biovalidation
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                                                                                                                                                                                                                                                                      Lehman
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Best Loc Matches

Local

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90.0%; 1.7%;

Score 16.8; D Pred. No. 1.6e 0; Mismatches

1.6e+03

DB 1;

Length

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Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ss; hybridisation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tion; capture oligonucleotide; pattern; mucosal; hair root;
germ cell; food additive; food supplement.
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28-FEB-2002; 2002DE-01008794 28-FEB-2002; 2002DE-01008794

(DEGS ) DEGUSSA BIOACTIVES GMBH Dieck HT, норре н;

WPI; 2003-714082/68

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable

Example; Page 4; 8pp; German.

cc reading out, where the nucleic acids are selectively bound using capture ce agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at cell the comprises all possible combinations of up to 10 nucleotides to binding of particular sorts of single stranded nucleic acids. The capture oligonucleotides that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LUA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by citation for the capture or agent and the anchor region conducting properties and especially in the form of a chip. Its surface by conducting properties and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface by stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular siaple stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the capture probes used in the matrix and phinding of single capture sieve. The method provides rapid, inexpensive and reproducible capture is a conditional compounds, e.g. food activity of pharmaceuticals and/or nutritional compounds, e.g. food capture single of the complete pattern of all nucleic acids from cells. It captures and the complete pattern of all nucleic acids from cells. It captures and compounds and compounds and compounds and compounds and cell, and capture probes used in the method of the invention. This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools,

BP; A, 0 ç; 0 G; 18 Η. 0 ς; 0

밁 cc comprise variable and non-variable regions, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture CC agents are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The material and/or membrane, having semication treath, gel, crystalline material and/or membrane, having semication treath, gel, crystalline material and/or membrane, having semication treath, gel, crystalline material and/or membrane, having semication treath and a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasilocovalent, supramolecular, cphysical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, capturity of pharmaceuticals and/or nutritional compounds, e.g. food cativity of pharmaceuticals and/or nutritional compounds. E.g. food cativity of pharmaceuticals and/or nutritional compounds, e.g. food cativity of pharmaceuticals and/or nutritional compounds. E.g. food cativity of pharmaceuticals and/or nutritional compounds. It can detect the method provides rapid, inexpensive and reproducible can detect very small differences in the nucleic acid poo RESULT 1578 Best Loca Matches Query Match Best Local This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides bears as a serious and the surface of a solid matrix and the surface of a solid matrix and (b) are a serious s Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable 04-SEP-2003. ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement. Rat DNA microarray capture oligonucleotide #4. 06-MAY-2004 (first entry) ADK01284 standard; 28-FEB-2002; 2002DE-01008794 28-FEB-2002; 2002DE-01008794. DE10208794-A1 Rattus sp. Example; Page 4; (DEGS ) DEGUSSA BIOACTIVES GMBH constant regions 428 TTTTATTTTATTTTTTTAA 447 18; Conservative μ TTTTTTTTTTTTTTAA 20 Dieck 8pp; German. DNA; HT, 90.0%; 21 норре н; ВP 0 Score 16.8; DB 1; Pred. No. 1.6e+03; Mismatches DB 1; Length 21; ٥, Gaps 0

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RESULT 1579
ADK01341
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CC comprise variable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region CC binding of particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The CC capture oligonucleotides are biotinylated and immobilised on a surface by comprises a sequence of 10-50, particularly 15-25, T residues. The CC interaction with strepcavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-CC is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, CC physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and cmixtures. The method provides rapid, inexpensive and reproducible of compresentation of differences in pools of nucleic acids from cells. It
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                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
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Pred. No. 1.
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RESULT 1580
ADK01283
ID ADK0128
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Matches
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agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food
                                                                                                                                                                                                                                    This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             allows imaging of the complete pattern of all nucleic acid in a cell, a can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent
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cell; food additive; food supplement.
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Pred. No. 1.6e
0; Mismatches
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additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, an can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention.
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Sequence 21 BP; 2 A; 1 C; 0 G; 18 T; 0 U; 0 Other;

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            Query Match
Best Local S
Matches 18
                   Local Similarity
 428
             18;
TTTTATTTTATTTTTTTAA 447
             Conservative
                  90.0%;
             0;
                   Score 16.8;
Pred. No. 1.
              Mismatches
                    .6e+03
                         DB 1; Length 21;
               Indels
               0
               Gaps
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0

RESULT 1581 ADK01331 片  $\vdash$ ΒP

ADK01331 standard; DNA; 21 ADK01331;

06-MAY-2004

(first entry)

Rat DNA microarray capture oligonucleotide #51.

ss; hybridisation; nerve; germ capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement.

Rattus

DE10208794-A1

04-SEP-2003

28-FEB-2002; 2002DE-01008794

28-FEB-2002; 2002DE-01008794.

DEGUSSA BIOACTIVES GMBH

Dieck HT,

Hoppe H;

WPI; 2003-714082/68 Boekenkamp D,

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable constant regions

Example; Page 5; 8pp; German

This invention describes a novel method for sorting single-stranded cc nucleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture captures that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides capture a 5'-invariable anchor region, the complement of which is present at comprise once in each nucleic acid and a 3'-variable, discriminatory region complement of the comprises all possible combinations of up to 10 nucleotides to allow that comprises all possible combinations of up to 10 nucleotides to allow comprises a sequence of single stranded nucleic acids. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The comprises a sequence of 10-50, particularly 15-25, T residues. The comprises a sequence of 10-50, particularly 15-25, T residues. The comprises a sequence of 10-50, particularly is of plastic, ceramic, glass, complete the surface by conducting properties and especially in the form of a chip. Its surface by conducting properties and especially in the form of and binding of single creaming and allowed and allowed mucleic acids. The surface is particularly a layer of (bio)molecular filaments and binding of single constants.

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additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, an can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food
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Sequence 21 BP; 1 A; 1 C; 0 G; 19 T; 0 U; 0 Other;

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Query Match
Best Local 9
Matches
18;
        Similarity
Conservative
        90.0%;
                1.7%;
0
        Score 16.8;
Pred. No. 1
 Mismatches
        .6e+03
                DB 1; Length 21;
 0
Gaps
  0
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밁 427 TTTTTATTTTATTTTTTTT 446 1 TTTTTTTTTTTTTTTTTA 20

ADK01330 standard; DNA; 21 ВP

ADK01330;

06-MAY-2004 (first entry)

Rat DNA microarray capture oligonucleotide

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus gp.

DE10208794-A1.

04-SEP-2003

28-FEB-2002; 2002DE-01008794

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH. Hoppe H;

WPI; 2003-714082/68

Boekenkamp D,

Dieck

HT,

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable constant regions

Example; Page 5; 8pp; German

RESULT 1582
ADK01330
XX
ADK0133
ADK0133
XX
ADK0133
A nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, This invention describes a novel method for sorting single-stranded

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RESULT 1583
ADK01332
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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                    Sorting single-stranded nucleic acid, patterns and screening active agents,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Boekenkamp D,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-MAY-2004
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                                                                                                                                                                                                                                                                                                                               Example;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DEGUSSA BIOACTIVES
                                                                                                                                                                                                                                                                                                                         Page 5; 8pp;
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cell; food additive; food supplement.
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                                                                                                                                                                                                                                                                                                                               German
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                    useful for analyzing expression uses capture agent with variable
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Novel isolated LPDL or LPDLR lipase polypeptides, useful for identifying substances that bind to the protein and which are useful for treating diseases associated with lipase function e.g. atherosclerosis and

hypercholesterolemia.

21-DEC-2001; 10-JAN-2002;

2001US-0341786P. 2002US-0346603P.

STEW/) (WENX/)

WEN X

STEWART A K.
TSUI L.
HEGELE R A.

Stewart AK,

Tsui ۲ 23-DEC-2002; 2002WO-CA001998.

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RESULT 1584
ADI2373
ID ADI2373
XX :
AC ADI2373
XX | Ipase;
KW | Iipase;
KW | Iipopro
XX | Iipopr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              lipase; LPDL; LPDLR; lipase deficiency; atherosclerosis; fatty liver disease; dyslipidaemia; hypercholesterolaemia; hypertriglyceridaemia; mixed dyslipidaemia; lipid deficient state; lipoprotein deficient state; human; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human LPDLR PCR primer #19
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18; Conserv
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Pred. No. 1.6e+03
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